Access DB# 66194

SEARCH REQUEST FORM

Scientific and Technical Information Center

Requester's Full Name:	,	Examiner #:	Date: '		
Art Unit: Phone No Mail Box and Bldg/Room Location:	umber 30	Serial Number:			
Mail Box and Bldg/Room Location:	R	esults Format Preferred	(circle): PAPER	DISK	E-N
lf mara than and soarch is submit	tted inlease nrio	ritize searches in order	of need.	:	
If more than one search is submit	*****	******	******	****	****
Please provide a detailed statement of the so Include the elected species or structures, ke utility of the invention. Define any terms the known. Please attach a copy of the cover sh	ywords, synonyms, a hat may have a specia	cronyms, and registry number I meaning. Give examples or	relevant citations,	n the coi	псері
Title of Invention:					
Inventors (please provide full names):					
Earliest Priority Filing Date:					
For Sequence Searches Only Please includ appropriate serial number.	e all pertinent informat	ion (parent, child, divisional, or	issued patent numbe	ers) along	with
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	Jan D Reference				
	Biotechnology &	703-308-4498	· ·	İ	
	jan.delava	l@uspto.gov			
STAFF USE ONLY	Type of Search	**************************************	d cost where appli	cable =	
Searcher:	NA Sequence (#)	STN	<u>: : : : : : : : : : : : : : : : : : : </u>		
Searcher Phone #: 4498	AA Sequence (#)	Dialog			
Searcher Location:	Structure (#)	Questel/Orbit	<u>:</u>		
Date Searcher Picked Up: 2\8 03	Bibliographic	Dr.Link		:	
Date Completed: 3-14 3	Litigation	Lexis/Nexis			
Searcher Prep & Review Time:	Fulltext	Sequence Systems			
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Clerical Prep Time: 30	Patent Family	www/Internet)			

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Database
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Perfect score:
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pir4:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	6	5	4	ω	2	1	Result No.
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toxin sensitivity	probable membrane	F6F9.11 protein -	DNA-directed RNA p	myosin II heavy ch	myosin heavy chain	chloride channel p	chloride channel p	transcription fact	protein T24E12.1 [probable ribosomal	protein nhr-58 (im	hypothetical prote	esterase/lipase l	degV protein homol	conserved hypothet	ribonuclease P (EC		рĭ	SOK2 protein - yea	transcription anti	C)	ש	hypothetical prote	Ω	-	Ω,		Ö	Description

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1 KQKIKHVVKLK

ALIGNMENTS

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C:Superfamily: hyaluronan receptor C:Keywords: glycoprotein; receptor F:260-382/Region: 21 residue repeats F:516-574/Region: hyaluronan binding #status predicted F:575-625/Region: hyaluronan binding #status predicted F:116,283,304,325,346,367,398,438,619/Binding site: car
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Species: Nus musculus (house mouse)
C;Date: 16-Nov-1995 #sequence_revision 08-Feb-1996 #text_change 01-Dec-2000
C;Accession: JC4298; A42925; A41923; S21586
R;Entwistle, J.; Zhang, S.; Yang, B.; Wong, C.; Li, Q.; Hall, C.L.; A, J.; Mowat, M.; Gene 163, 233-238, 1995
A;Title: Characterization of the murine gene encoding the hyaluronan receptor RHAMM.
A;Reference number: JC4298; MUID:96011639; PMID:7590272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Note: authors translated the codon AGT for residue 377 as Thr and CTA for residue 5 R;Hardwick, C.; Hoare, K.; Owens, R.; Hohn, H.P.; Hook, M.; Moore, D.; Cripps, V.; Au J. Cell Biol. 117, 1343-1350, 1992
A;Title: Molecular cloning of a novel hyaluronan receptor that mediates tumor cell mo A;Reference number: A41923; MUID:92299690; PMID:1376732
A;Accession: A41923
                                                                                                                                                                                                                                                                                                                                                                                   A; Note: this sequence has been corrected in reference A42925 C; Comment: This protein regulates cell motility and transformation,
                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule type: mRNA
A;Residues: 155-376,'S',378-504,'E',506-507,'L',508-630 <HA2>
A;Cross-references: GB:X64550
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A; Residues: 155-376,'S',378-504,'E',506-631 <HAR>
A; Cross-references: GB:X64550
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A;Contents: erratum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: mRNA
A; Residues: 1-631 <ENT>
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                                                                                                                                                                                                                                                                                                A; Introns: 20/3; 54/2; 79/2; 137/3; 187/3; 259/2; 382/2; 431/2; 482/2; 515/3; 574/3;
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                                 Similarity
100.0%; ilarity 100.0%; Conservative 0
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Pred. No. 0.054;
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C;Species: Now Sapiens (man)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 29-Sep
C;Accession: JC5016
R;Wang, C; Entwistle, J:; Hou, G.; Li, O.; Turley, E.A.
Gene 174, 299-306, 1996
A;Title: The characterization of a human RHAMM cDNA: Conservation of t
A;Reference number: JC5016; MUID:97045829; PMID:8890751
A;Contents: breast
A;Accession: JC5016
A;Molecule type: mRNA
A;Residues: 1-725 <WAN>
A;Cross-references: GB:U29343
A;Note: it is uncertain whether Met-1 or Met-196 is the initiator
C;Comment: This receptor regulates focal adhesion turnover, and regulations of the conservations:
A;Gene: GDB:HMMR; RHAMM
A:Cross-references: CDR:A87204: OMTM-600476
hypothetical protein SA2433 [imported] - Staphylococcus aureus (strain N315) C:Species: Staphylococcus aureus C:Species: Staphylococcus aureus C:Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 22-Oct-2001 C:Accession: H90071 R:Gobayashi, I: Baba, T: Yuzawa, H: Kobayashi, I: Cui, L: ma, A.; Mizutani-Ui, Y.; Kobayashi, N.; Sawano, T: Inoue, R: Kaito, C: Sekimizu, C: Shiba, T: Hattori, M: Ogasawara, N.; Hayashi, H.; Hiramatsu, K. Lancet 357, 1225-1240, 2001
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H90071
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A; Note: T30N20_260
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A; Residues: 1-148 <BEV>
A; Cross-references: EMB
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A; Accession: T50808
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C:Specles: Homo sapiens (man)
C:Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change
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C; Superfamily: )
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se: cultivar Columbia;
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3; Mismatches
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hypothetical protein F8F6.230 - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #
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C;Superfamily: yeast DNA-directed
C;Keywords: nucleomorph
                                                  A; Note:
                                                                A; Map position: 5
A; Introns: 100/3
                                                                                                                                               A; Molecule type: DNA
A; Residues: 1-1495 <BEV>
                                                                                                                                                                                                A;Reference number: Z24488
A;Accession: T48429
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R;Bevan, M.; Peters, S.A.;
submitted to the Protein S
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A: Residues: 1-624 <KUR>
A: Cross-references: GB:BA000018; PID:g13702598;
A: Cross-references: Strain N315
                                                                                                               A; Experimental source:
                                                                                                                                  A; Cross-references:
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C;Genetics:
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A; Residues: 1-1473 < DOU>
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R;Douglas, S.; Zauner, S.; Fraunholz, M.;
Nature 410, 1091-1096, 2001
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                                                  . 230
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                             EMBL:AL162873
be: cultivar Columbia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GB:AJ010592;
                                                                                                                                                                                                                             Sequence
   66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            68.5%;
70.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   68.5%;
                                                                                                                                                                                                                                                van
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genome of an enslaved algal nucleus MUID:11323671; PMID:11323671
                                                                                                                                                                                                                                              Staveren,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ω,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NID: g12580785;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 37;
Pred. No.
 Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2
                                                                                                                                                                                                                               Database,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 37;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RNA polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                            20-Apr-2000 #text_change 20-Apr-2000
37;
No.
                                                                                                                                                                                                                                              X
                                                                                                               BAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Beaton,
 DB 2;
1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB
66;
                                                                                                                                                                                                                                March
                                                                                                                                                                                                                             Dirkse, W.;
March 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 2;
1.6e+02;
                                                                                                               clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PIDN:CAC27103.1; GSPDB:GN00151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PIDN: BAB43738
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            III
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>ب</u>
               Length 1495;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 1473;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            160K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                              Stiekema,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GSPDB: GN00149
                                                                                                                                                                                                                                            W.; Bancroft,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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Similarity

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C;Genetics:
A;Gene: VC1678
A;Map position:
                                                                                                                                                                                                                                                                                            Science 294, 849-852, 2001

A; Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; A; Title: Comparative genomics of Listeria species.

A; Reference number: AB1077; MUID:21337279; PMID:11679669

A; Accession: AB1112
A; Accession: AB1112
A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-623 <GLA>
A; Cross-references: GB:NC_003210; PIDN:CAD00824.1; PID:g16409662; GSPE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Species: Listeria monocytogenes
C;Date: 27-Nov-2001 #sequence_revision 27-Nov-2001
C;Accession: ABIL12
R;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend,
Dominguez-Bernal, G.; Duchaud, E.; Durand, L.;
                                                                                                                          Qy
                                                                                                                                                                                                                                                         A;Experimental source: strain EGD-e
C;Genetics:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-222 <HEI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Title: DNA Sequence of both chromosomes of the cholera pathogen Vibrio cholerae A;Reference number: A82035; MUID:20406833; PMID:10952301 A;Accession: H82169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Heidelberg, J.F.; Eisen, J.A.; Nelson, W.C.; Clayton, R.A.; Gwinn, M.L.; chardson, D.; Ermolaeva, M.D.; Vamathevan, J.; Bass, S.; Qin, H.; Dragoi, Il, R.R.; McKalanos, J.J.; Venter, J.C.; Fraser, C.M.
Nature 406, 477-483, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           phage shock protein A VC1678 [imported] - Vibrio cholerae (strain N16961 serogroup O1)
C;Species: Vibrio cholerae
C;Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 02-Feb-2001
C;Accession: H82169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Qγ
                                                                                           В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 В
transcription antiterminator (BglG
                   AF1473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       D.; Jones, L.M.; Karst,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Experimental source: serogroup
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Cross-references: GB: AE004246;
                                     RESULT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transcription antiterminator (BglG family) homolog lmo0297 [imported] - Listeria monocy
                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                              QKIKHVVKLK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KQKIKHVVK 9
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                                                                                                                                                                   Similarity 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
6; Conser
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                                                                                                                                                                   Conservative
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66.7%;
                                                                                                                                                                                     66.7%;
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Pred.
3; Mis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ψ
                                                                                                                                                                                 Score 36;
Pred. No.
family) homolog lin0325 [imported]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-Nov-2001 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      35;
                                                                                                                                                                                   DB 2;
1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , A.; Baquero, F
Dussurget, O.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                      Length 623;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 222
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El Tor
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                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                   Gaps
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 1
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Voss, H.; We
Listeria innocua
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                                                                                                                                                                   0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bloecker
Fsihi, H.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Wehland
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Α.,
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                                 A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-835 <BEV>
                                                                                         A; Reference number: Z24489
A; Accession: T48260
                                                                                                                                        C;Accession: T48260
R;Bevan, M.; Hilbert, H.; Braun, M.; Holzer,
                                                                                                                                                                                 hypothetical protein T1E22.150 - Arabidopsis thallana C;Species: Arabidopsis thallana (mouse-ear cress) C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 20-Apr-2000
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A; Experimental source: cultivar Columbia;
               A; Cross-references:
                                                                                                                            submitted to the Protein Sequence Database,
                                                                                                                                                                                                                                           T48260
                                                                                                                                                                                                                                                             RESULT 11
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EMBL: AL162874

BAC

clone

T1E22

March E.; Brandt,

2000

A.; Duesterhoeft, A.;

Bancro

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A; Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; A; Title: Comparative genomics of Listeria species.
A; Reference number: AB1077; MUID:21537279; PMID:11679669
A; Accession: AF1473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .; Dominguez-Bernal, G.; Du D.; Jones, L.M.; Karst, U. Science 294, 849-852, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C; Species: Listeria
C; Date: 27-Nov-2001
C; Accession: AF1473
                                                                                                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-785 <LYE>
A; Cross-references: EMB
                                                                                                                                                                                                                                                                                                                                              A; Reference number: S54014
A; Accession: S54016
                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOK2 protein - yeast (Saccharomyces cerevisiae)
N;Alternate names: protein YM9711.03c; protein YMR016c
C;Specles: Saccharomyces cerevisiae
C;Date: 08-Jul-1995 #sequence_revision 01-Sep-1995 #text_change 06-Feb-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Gene: lin0325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: GB:AL592022; PIDN:CAC95558.1; A;Experimental source: strain Clip11262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-623 <GLA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.; Baquero, F.; Berche, P.; Bloec.; Dominguez-Bernal, G.; Duchaud, E.; Durand, L.; Dussurget, O.; Entian, K.D.; Fsihi,
                                                                                                                                                                                              A; Cross-references:
                                                                                                                                                                                                                A; Gene: SGD: SOK2
                                                                                                                                                                                                                                                           A; Experimental source: strain
                                                                                                                                                                                                                                                                                                                                                                                           submitted to the EMBL Data
                                                                                                                                                                                                                                                                                                                                                                                                                                 C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Status: preliminary
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Species: Listeria innocua; Date: 27-Nov-2001 #text_change; Date: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change
                                                                                                                           Query Match
                                                                                                                                                                                                                                         Genetics:
                                                                                     Matches
469 EKIRHVVKI 477
                                                                                                                                                                        position:
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                                                                                     Local Similarity
nes 6; Conser
                                          2 QKIKHVVKL 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                   S54016
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                                                                                     Conservative
                                                                                                                                                                                                                                                             EMBL:Z49211; NID:g798922; PID:g798925; pe: strain AB972
                                                                                                                                                                                              SGD:S0004618;
                                                                                                         66.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 36; DB
Pred. No. 1e+C
3; Mismatches
                                                                                   Score 36; DB
Pred. No. 1.3e
3; Mismatches
                                                                                                                                                                                              MIPS:YMR016c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 2;
1e+02;
                                                                                                       .3e+02;
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                                                                                                                           Length 785;
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                                                                                     0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Voss, H.;
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Wehla
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RESULT 14
B89836
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C:Species: Xenopus laevis (African clawed frog)
C:Date: 22-Oct-1999 #sequence_revision 22-Oct-1999 #text_change 02-Jun-2000
C:Accession: T30335
R:Boleti, H.; Karsenti, E.; Vernos, I.
Cell 84, 49-59, 1996
A:Title: Xklp2, a new Xenopus centrosomal kinesin-like protein required for A;Reference number: Z20827; MUID:96140639; PMID:8548825
A;Accession: T30335
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-1388 <BOL>
A;Cross-references: EMBL:X94082; NID:g1129172; PID:e213754; PIDN:CAA63826.1
                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Notacus, France, DNA
A; Molecule type: DNA
A; Residues: 1-117 <STO>
A; Cross-references: GB:AE005176; PID:g12722976; PIDN:AAK04226.1; GSPDB:GN00146
                                                                                                                                    Вþ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R; Bolotin, A.; Wincker, P.; Mauger, S.; Jaillon, O.; Malarme, Genome Res. 11, 731\text{-}753, 2001
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C;Superfamily: unassigned kinesin-related proteins; kinesin motor
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C;Genetics:
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A; Map position: 5
A; Introns: 53/1; 82/3;
A; Note: T1E22.150
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degv protein homolog K05_orf237 - Mycoplasma pneumoniae (strain ATCC 29342)
C;Species: Mycoplasma pneumoniae
A;Variety: ATCC 29342
C;Date: 27-Feb-1997 #sequence_revision 25-Apr-1997 #text_change 07-Dec-1999
C;Accession: S73504
                                                                                                 A;Note: the nucleotide sequence C;Genetics:
                                                                                                                                                                                                                                  Nucleic Acids Res. 24, 4420-4449, 1996
A;Title: Complete sequence analysis of the genome of the A;Reference number: S73327; MUID:97105885; PMID:8948633
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C; Accession: B89836
R; Kuroda, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.;
Ra, A.; Mizutani-Ul. Y.; Kobayashi, N.; Sawano, T.; Inoue, R.; Kaito, C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K.
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                                                                                 A; Genetic
                                                                                                                                          A; Cross-references:
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A; Residues: 1-237 <HIM>
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A; Residues: 1-168 <KUR>
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A;Title: Whole genome sequencing of meticillin-resistant A;Réference number: A89758; MUID:21311952; PMID:11418146
                                                                                                                                                                                           A; Status: preliminary; nucleic acid sequence not shown; translation
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Hyaluronan mediated motility receptor (Intracellular hyaluronic acid binding protein) (Receptor for hyaluronan-mediated motility) (CD168
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ERK REGULATION, AND SUBCELLULAR LOCATION MEDLINE-98225222; PubMed-9556628; Zhang S., Chang M.C., Zylka D., Turley S. "The hyaluronan receptor RHAMM regulates
                                                   MEDLINE=94308286; PubMed=7518470;
Hall C.L., Wang C., Lange L.A., Turley E.A.;
"Hyaluronan and the hyaluronan receptor RHAMI
turnover and transient tyrosine kinase activ.
J. Cell Biol. 126:575-588(1994).
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"Molecular cloning of
cell motility.";
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Hyaluronan mediated motility receptor (Intracellular hyaluronic acid binding protein) (Receptor for hyaluronan-mediated motility).
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Mowat M., Greenberg A.H.,
"Characterization of the
RHAMM.":
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Mammalia; Eutheria; Rodentia;
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                                                                                                                             Cell Biol.
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                                                                                                                            117:1343-1350(1992).
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Sciurognathi; Muridae;
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Ponta H., Herrlich P.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CARBOHYD
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J. Biol. Chem.
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AF079222; AAD08670.1;
AL; AJ005919; CAA06768.1;
AJ005921; CAA06768.1;
AJ005921; CAA06768.1;
AJ005922; CAA06768.1;
AJ005923; CAA06768.1;
AJ005923; CAA06768.1;
LL; AJ005923; CAA06768.1;
LL; AJ005923; CAA06768.1;
LL; AJ005923; CAA06768.1;
LL; AJ005923; CAA06848.1;
AG450; CAA45849.1;
AG450; CAA45848.1;
MGI:104667; Hmmrr.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EXTRACELLULAR-REGULATED KINASE (ERK) ACTIVITY.
SUBUNIT: SUBUNIT OF THE HARC COMPLEX.
SUBCELLULAR LOCATION: CELL SCHOOL CYTOPLASMIC.
ALTERNATIVE PRODUCTS: 2 ISOFORMS; RHAMM1V4 (SHOWN HERE) AND RHAMM1; ARE PRODUCED BY ALTERNATIVE SPLICING.
TISSUE SPECIFICITY: UBIQUITOUSLY EXPRESSED.
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-i- FUNCTION: PLAYS A GENERAL REGULATORY ROLE IN THE CYCLIC AMP-DEPENDENT PROTEIN KINASE-STIMULATED (PKA) SIGNAL TRANSDUCTION PATHWAY BY REGULATING THE EXPRESSION OF GENES IMPORTANT IN GROAND DEVELOPMENT. MAY INHIBIT THE SWITCH FROM UNICELLULAR TO AND DEVELOPMENT. MAY INHIBIT THE SWITCH FROM UNICELLULAR TO
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SOK2 OR YMR016C OR YM9711.03C
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the European Bioinformatics Institute.
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                                                                                                                                                                                                SEQUENCE
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MEDLINE=96069395; PubMed=8524252;

Ward M.P., Gimeno C.J., Fink G.R., Garrett S.;

Ward M.P., Gimeno C.J., Eink G.R., Garrett S.;

"SOK2 may regulate cyclic AMP-dependent protein kinase-stimulated growth and pseudohyphal development by repressing transcription.";

Mol. Cell. Biol. 15:6854-6863(1995).
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SIMILARITY: BELONGS TO THE EFG1/PHD1/STUA FAMILY.
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Last annotation updat
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POLY-GLN.
POLY-TYR.
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                                                                                             Score 36; DB
Pred. No. 40;
3; Mismatches
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Matches
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Best Local
                                                                             01-NOV-1997 (Rel. 35, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
Hypothetical protein MG450 homolog (K05_orf2 MPN664 OR MP178.
                                                                                                                                                                 Y450_MYCPN
P75127;
                                                          Mycoplasma pneumoniae Bacteria; Firmicutes;
                                                                                                                                                                                                                                                                                                                                                                                     Complete
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboratic between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use by non-profit institutions as long as its content is in no was modified and this statement is not removed. Usage by and for commercia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lactis ssp. lactis IL1403.";
Genome Res. 11:731-753(2001)
-!- FUNCTION: RNaseP catalyz
                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR000100; Ribonuclease_P.
Pfam; PF00825; Ribonuclease_P; 1.
TIGREAMS; TIGR00188; rnpA; 1.
PROSITE; PS00648; RIBONUCLEASE_P; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09CJ73:
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Ribonuclease p protein component (EC 3.1.26.5) (RNaseP protein) (RNase
STRAIN-ATCC 29342 /
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; AE006251; AAK04226.1; -. HSSP; P25814; 1A6F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE~21235186; PubMed=11337471;
BOlotin A., Wincker P., Mauger S., Jaillon O.,
Welssenbach J., Ehrlich S.D., Sorokin A.;
"The complete genome sequence of the lactic ac.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. STRAIN-IL1403;
               SEQUENCE FROM
                                            NCBI_TaxID=2104;
                                                                                                                                                                                                                                                                                                                                                                                                                Hydrolase; Nuclease; Endonuclease; tRNA processing; RNA-binding;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or send
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FUNCTION: RNASSP Catalyzes the removal of the 5'-leader sequence from pre-trnA to produce the mature 5'terminus. It can also cleave other RNA substrates such as 4.5s RNA. The protein component plays an auxiliary but essential role in vivo by binding to the 5'-leader sequence and broadening the substrate specificity of the ribozyme (By similarity).

CATALYTIC ACTIVITY: Endonucleolytic cleavage of RNA, removing 5'-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein subunit (By similarity).
SIMILARITY: BELONGS TO THE RNPA FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        extra-nucleotide from tRNA precursor.
SUBUNIT: Consists of a catalytic RNA component
                                                                                                                                                                                                                                                                KKNLKHVLKL 107
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                                                                                                                                                                                                                                                                                                                         Similarity
6; Conser
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117 AA;
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illarity 60.0%;
Conservative
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es; Lactobacillales; Streptococcaceae;
M129
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                                                            Mollicutes;
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                                                          Mycoplasmataceae; Mycoplasma
                                                                                                     ion update)
(K05_orf237).
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Best Local
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                                                                                                                                                                                                                                                     rRNA gene of Fehrchard.";
other filamentous fungi.";
J. Biochem. 117:888-896(1995).
-I- FUNCTION: ESSENTIAL FOR MITOCHONDRIAL PROTEIN S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-FEB-1996 (Rel. 33, Created)
01-FEB-1996 (Rel. 33, Last seque
01-NOV-1997 (Rel. 35, Last annot
Mitochondrial ribosomal protein
                                             SEQUENCE
                                                                                                             use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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01-FEB-1996
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InterPro; IRR003797; DegV.
Pfam; PF02645; DUF194; 1.
Hypothetical protein; Complete
SEQUENCE 237 AA; 26839 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                         EMBL; D14567; BAA03421.1; -.
                                                                                                                                                                                                                                                                                                                                                             STRAIN-NRRL 2159A; MEDLINE-96070189; PubMed-759255;
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                                                                                                                                                                                                                                                                                                                 Yamamoto H., Naruse A., Ohsaki T., Sekiguo
"Nucleotide sequence and characterization
"RNA gene of Penicillium urticae, and its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic Acids Res.
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                                                                                                                                                                                                                   SIMILARITY: BELONGS TO THE VAR1 FAMILY OF RIBOSOMAL PROTEINS
                                                                                                                                                                                                                                 (BY SIMILARITY). SUBCELLULAR LOCATION: Mitochondrial
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                                                                                                    an email to license@isb-sib.ch).
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                                                         protein;
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                               Mitochondrion.
Mitochondrion.
M7447 MW; 38A8263B45BB4041 CRC64;
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54.5%;
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DOMAIN
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Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes; Saccharomycetales; mitosporic Saccharomycetales; Candida.
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01-NOV-1995 (Rel.
30-MAY-2000 (Rel.
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           Chloride
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Pfam; PF02292; Yeast_DNA_bind; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             PSEUDOHYPHAL MORPHOGENESIS.
-!- SUBCELLULAR LOCATION: Nuclear (Potential).
-!- SIMILARITY: BELONGS TO THE EFG1/PHD1/STUA FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           regulating morphogenetic processes in fungi."; EMBO J. 16:1982-1991(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Candida
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN-ATCC
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                                                                                                                                                                Similarity 5; Conser
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protein 7 (CIC-7).
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CLC7_RAT STANDARD; PRT; 8
ID CLC7_RAT STANDARD; PRT; 8
AC p51799;
AC p51799; (Rel. 34, Created)
DT 01-OCT-1996 (Rel. 34, Last sequence up
DT 101-UN-2002 (Rel. 41, Last annotation
DE Chloride channel protein 7 (ClC-7).
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Pfam; PF00654; voltage_CLC;
PRINTS; PR00762; CLCHANNEL.
SMART; SM00116; CBS; 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; AF063101; AAC18832.1;
EMBL; AF063098; AAC18832.1;
EMBL; AF063099; AAC18832.1;
EMBL; AF063100; AAC18832.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Cell 104:205-215(2001)
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Friedrich W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
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                                                                                                                            215 KIPHVVRLK 223
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SIMILARITY: CONTAINS 2 CBS DOMAINS.
                                                                                                                                                                                                                                                                                                                                                                   domain;
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Kasper D., Bosl M.R., Kaiser E.,
W., Delling G., Jentsch T.J.;
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Best Local :
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               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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NCBI_TaxID=10116;
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InterPro: IPR001644: CBS_domain.
InterPro: IPR001807; C1-channel_volt.
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"C1C-6 and C1C-7 are two novel broadly expressed chloride channel family.";
FEBS Lett. 377:15-20(1995).
NCBI_TaxID=9606;
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Pfam; PF00654; voltage_CLC;
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                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
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                                                                                                                                                                                                                                                       215
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                                                                                                                                                                                                                                                                                      3 KIKHVVKLK 11
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SUBCELLULAR LOCATION: Integral membrane protein.
TISSUE SPECIFICITY: BRAIN, TESTIS, MUSCLE AND KIDNEY.
SIMILARITY: BELONGS TO THE CHLORIDE CHANNEL FAMILY.
SIMILARITY: CONTAINS 2 CBS DOMAINS.
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CBS 2.
; A7D6AA3D21DA6FED CRC64;
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EMBL; BC012737; AAH12737.1; -.
EMBL; BC743, CAA91556.1; -.
EMBL; AJ001910; CAA05083.1; -.
EMBL; U88844; AAB48530.1; -.
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MIM; 602727;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Schutte B.C., Malik M.I., Fingert J., Stone E., Lamb F.S.;
Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases.
-i- FUNCTION: VOLTAGE-GATED CHLORIDE CHANNEL. CHLORIDE CHANNELS
SEVERAL FUNCTIONS INCLUDING THE REGULATION OF CELL VOLUME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Brandt S., Jentsch T.J.;
"ClC-6 and ClC-7 are two novel broadly expressed chloride channel family.";
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InterPro; IPR001807; C1-channel_volt.
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TISSUE-Breast;
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"The exon-intron architecture of human chloride
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SUBCELLULAR LOCATION: Integral membrane protein.
TISSUE SPECIFICITY: BRAIN, TESTIS, MUSCLE AND KI
SIMILARITY: BELONGS TO THE CHLORIDE CHANNEL FAM.
SIMILARITY: CONTAINS 2 CBS DOMAINS.
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n channels in
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98234303; PubMed=9565675;
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Best Local
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                                                   Pfam; PF01576; Myosin_tail; 1.
Pfam; PF02736; Myosin_N; 1.
PRINTS; PR00193; MYOSINHEAVY.
ProDom; PD000355; myosin_head; 1
SMART; SM00015; IQ; 1.
SMART; SM00042; MYSC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CONFLICT
CONFLICT
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                          01-JUN-1994
01-JUN-1994
16-OCT-2001
                                                                                                                                                                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through a collab
between the Swiss Institute of Bioinformatics and the EMBL outst
the European Bioinformatics Institute. There are no restrictions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilidae; Drosophila. NCBI_TaxID=7227;
                                                                                                                                                                             FlyBase; FBgn0005634; zip. InterPro; IPR000048; IQ_region.
                                                                                                                                                                                                      HSSP;
                                                                                                                                                                                                                            PIR;
                                                                                                                                                                                                                                                           entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                 modified and this statement
                                                                                                                                                                                                                                                                                                                                                 -i- ALTERNATIVE PRODUCTS: 2 ISOFORMS; A LONG FORM (SHOWN HERE) AND SHORT FORM; ARE PRODUCED BY ALTERNATIVE SPLICING.
-i- SIMILARITY: CONTAINS 1 MYOSIN-LIKE GLOBULAR HEAD DOMAIN.
-i- SIMILARITY: CONTAINS 1 IQ DOMAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MYSN_DROME Q99323;
                      Myosin; Alternative splicing; Coiled coil; Actin-binding; ATP-binding; Calmodulin-binding.
                                             PROSITE;
                                                                                                                                            InterPro; IPR004009; Myosin_N.
InterPro; IPR002928; Myosin_tail.
InterPro; IPR001609; myosin_head.
                                                                                                                                                                                                                 PIR; B36014; B36014.
                                                                                                                                                                                                                                      EMBL; M35012; AAA28713.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE-Embryo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Myosin heavy
                                                                                                                                     Pfam;
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                                                                                                                                                                                                                            A36014; A36014.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KIKHVVKLK 11
                                                                                                                        PF00063; myosin_head; PF00612; IQ; 1.
                                                                                                                                                                                                     P10587; 1BR2
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7; Conserv
                                             PS50096; IQ;
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279
805
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(Rel. 29, Last sequence update)
(Rel. 40, Last annotation update)
chain, non-muscle (Zipper protein)
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279
88679
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77.8%;
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Pred. No.
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MÝOSIN HEAD-LIKE
IQ.
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-> L (IN REF. 3 AND 4).
E56BC0B4ADE1C695 CRC64;
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RESULT 13
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Best Local :
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P27625;
01-AUG-1992 (Rel. 2
01-AUG-1992 (Rel. 2
15-DEC-1998 (Rel. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDILINE PARIA DAY, FRANKA M., Gu H., Fox B.A., Inserburg J.,
Li W.B., Bark D.J., Tanaka M., Gu H., Fox B.A., Inserburg J.,
"Characterization of the gene encoding the largest subunit of
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NP_BIND
DOMAIN
Transferase: DNA-directed RNA polymerase;
Zinc-finger: Nuclear protein.
ZN_FING 88 101 C3H-TYPE (POT
SEQUENCE 2339 AA; 272829 MW; EDD899363
                                                                                     Pfam; PF00623; RNA_pol_A; 1 Pfam; PF01854; RNA_pol_A2; :
                                                                                                                                                                                                             or send
                                                                                                                                                                                                                                                       use by non-profit institutions as long modified and this statement is not removed.
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                                                                                                                                                                                                                                                                                                                                          This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=92018020; PubMed=1656254;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plasmodium falciparum.
Eukaryota; Alveolata; Apicomplexa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1063 EEKAKHLAKLK 1073
                                                                                                                        InterPro; IPR000722; RNA_pol_A.
InterPro; IPR002879; RNA_pol_A2
                                                                                                                                                                        EMBL; M73770; AAA29729.1; -
                                                                                                                                                                                                                                      entities requires a
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                                                                                                                                                                                                                                                                                                                                                                                PRECURSOR, POLYMERASE II FOR THE MRNA PRECURSOR, AND POLYMERASE III FOR 5S AND TRNA GENES.
SIMILARITY: BELONGS TO THE RNA POLYMERASE BETA' CHAIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                             SUBCELLULAR LOCATION: Nuclear.
MISCELLANEOUS: THREE DISTINCT ZINC-CONTAINING RNA POLYMERASES FOUND IN EUKARYOTIC NUCLEI: POLYMERASE I FOR THE RIBOSOMAL RNI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUBSTRATES.
CATALYTIC ACTIVITY: N nucleoside triphosphate = N diphosphate
                                                                                                                                                                                                                                                                                                 European Bioinformatics Institute.
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                                                                                                                                                                                                                                                                                                                SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation
                                                                                                                                                                                                                an email to license@isb-sib.ch).
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250
682
705
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STANDARD;
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54.5%;
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Last annotation update)
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MISSING (IN SHORT ISO
MW; 73E3CB02BA8F2528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COILED
ATP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REACTIVE SULFHYDRYL/ACTIN-BINDING LIGHT MEROMYOSIN (LMM).
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ACTIN-BINDING.
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1.6e+02;
2;
                          (POTENTIAL)
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      CRC64;
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Query

Match

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                                             Matches
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015258; 095322;
30-MAY-2000 (Rel. 3
30-MAY-2000 (Rel. 3
30-MAY-2000 (Rel. 3
                                                                                                                                          TRANSMEM
TRANSMEM
CONFLICT
                                                                                      SEQUENCE
                                                                                                                                                                                               Pfam;
                                                                                                                                                                                                                             EMBL;
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                                                                                                                                                                                                                                                                                                                                                                    Barrow I.K.-P., Boguski M.S., Touchman J., Spencer F.;
Submitted (AUG-1998) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: INVOLVED IN THE RETRIEVAL OF ENDOPLASMIC RETICULUM
MEMBRANE PROTEINS FROM THE EARLY GOLGI COMPARTMENT (BY
                                                                                                                                                                                                                     EMBL;
                                                                                                                                                                                                                                                              modified and this statement is not removed. entitles requires a license agreement (See )
                                                                                                                                                                                                                                                                                                                                     SIMILARITY).
-!- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN. GOLGI.
-!- SIMILARITY: BELONGS TO THE RER1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                     Transmembrane;
                                                                                                                                                                                                                                                   or send
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                                                                                                                                                                                                                                                                                                                                                                                                                                                cerevisiae.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fuellekrug J., Boehm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE-Breast cancer;
MEDLINE-97454982; Pub
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163
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 KRQIKHMIKYR
                     KOKIKHVVKLK
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AF091071; AAC72940.1;
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5; Conser
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5; Conserv
                                                                                                                                                                                                                                                    an email to license@isb-sib.ch).
                                                                                                                                                                                                       IPR004932; Rer1.
                                                                                      196
                                           Conservative
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160
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Primates;
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POTENTIAL.

HAATKGILVAMVCTFFDAFNVPVFWPILVMYFIMLFCITMK
RQIKHMIKYRYLPFTHGKRRYRGKEDAGKAFAS -> DASV
CGDGRCSCKAGGGRQCPVLAADAALTFSPHLKACGYQGHPC
                                                    Score 34; DB
Pred. No. 24;
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4; Mismatches
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Q1-NOV-1997 (Rel. 35, C
Q1-NOV-1997 (Rel. 35, I
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Puttative modification n
MJAIVMP OR MJ1328
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MEDLINE-96337999; PubMed=8688087;

Bult C.J., White O., Olsen G.J., Zhou L., Fleischmann R.D.,

Sutton G.G., Blake J.A., FitzGerald L.M., Clayton R.A., Gocayne J.D.,

Kerlavage A.R., Dougherty B.A., Tomb J.-F., Adams M.D., Reich C.I.,

Overbeek R., Kirkness E.F., Weinstock K.G., Merrick J.M., Glodek A.,

Scott J.L., Geoghagen N.S.M., Weidman J.F., Fuhrmann J.L., Nguyen D.,

Utterback T.R., Kelley J.M., Peterson J.D., Sadow P.M., Hanna M.C.,

Cotton M.D., Roberts K.M., Hurst M.A., Kaine B.P., Borodovsky M.,

Klenk H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

"Complete genome sequence of the methanogenic archaeon, Methanococcus langaschii."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hypothetical protein; Transferase; Methyltransferase; Restriction system; Complete proteome. SEQUENCE 298 AA; 35328 MW; 9BE52F35B001822B CRC64
                                                                                                                                                                                                                                                                                                                                                                             REBASE; 3895;
TIGR; MJ1328;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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completed: February me : 8.96667 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                s SWISS-PROT entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation - European Bioinformatics Institute. There are no restrictions on tist by non-profit institutions as long as its content is in no way iffed and this statement is not removed. Usage by and for commercial
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sp_fungi:*
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Q9UDA
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Q92026 mus sp. hya
Q9w1f7 rattus norv
Q8wdr2 mus musculu
Q9fhm8 arabidopsis
Q8s8k9 arabidopsis
P93142 forsythia i
Q9avx8 guillardia
Q9leu2 arabidopsis
Q9vbu6 drosophila
Q99qz8 staphylococ
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DUN-2002 (TrEMBLrel. 21, Last annotation update)
Hyaluronic acid-mediated motility receptor (Fragment)
Ovis aries (Sheep)
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Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF310973; AAK69578.1; -.
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BOVIdae; Caprinae; Ovis.
NCBI_TaxID=9940;
                                                                                                                                                                                                                             Aoki T.A., Forteza R.M., "Receptor for hyaluronic
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acid mediated motility in sheep
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SEQUENCE FROM N.A.
MEDLINE-92299690; PubMed-1376732;
Hardwick C., Hoare K., Owens R., Hohn H.P., Hook M.,
Hardwick C., Hoare K., Owens D.M., Turley E.A.;
Nance D.M., Turley E.A.;
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01-JUN-2002
01-JUN-2002
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SEQUENCE
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J. Cell Biol. 118.753-753(1992).
EMBL; S41029; AAA09809.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cripps V., Austen L., Nance D.M., Turley E.A.; "Molecular cloning of a novel hyaluronan receptor cell motility.";
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bovidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bos taurus (Bovine).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RHAMM
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                                                                                           401
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                                                                                                KOKIKHVVKLK 411
                                                                                                                                         KOKIKHVVKLK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KQKIKHVVKLK 38
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                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                   476 AA;
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(TrEMBLrel.
(TrEMBLrel.
r hyaluronic
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(TrEMBLrel. 19, Last sequence update)
(TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                          Conservative
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larity 100.0%;
Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
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Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9452 MW;
                                                                                                                                                                                                                                                                                   55486 MW;
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21, Last sequence update)
21, Last annotation update)
21, Last annotation (France)
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Pred. No. 0.0
); Mismatches
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                                                                                                                                                                                       Score 54; DB 11
Pred. No. 0.094;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Craniata; Vertebrata; Euteleosto
Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRT;
                                                                                                                                                                                                                                                                                   62DCA82045EE0CB2
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                                                                                                                                                                                                                                                                                   CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 82;
                                                                                                                                                                                                                                   Length
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                                                                                                                                                                                                                                                                                                                                                                                                   receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Moore
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REVENT.4
REV
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Q9FHM8
ID Q9
AC Q9
DT 01
DT 01
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Best Local S
Matches 11
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Q9FHM8;
01-MAR-2001
01-MAR-2001
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Q9WUF7;
01-NOV-1999
01-NOV-1999
01-JUN-2001
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Q8VDR2;
                                                                        Q9FHM8
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SEQUENCE
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(TrEMBLrel.
                                                                        PRELIMINARY;
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Query Match
Best Local Similarity
                                                                                                                                    Strausberg R.;
Submitted (JAN-2002)
EMBL; BC021427; AAH2;
                                                                                                                                                                                                                                                            01-MAR-2002 (TrEMBLrel. 20, Cre
01-MAR-2002 (TrEMBLrel. 20, Las
01-MAR-2002 (TrEMBLrel. 20, Las
5imilar to hyaluronan mediated
                                                                                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=PHEOCHROMOCYTOMA;
Lynn B.D., Li X., Cattini P.A., Nagy J.I.;
Lynn B.D., Li X., Cattini P.A., Nagy J.I.;
"Sequence, protein expression and erk associat
"RHAMM in PC12 cells.";
Submitted (JAN-2001) to the EMBL/GenBank/DDBJ
EMBL; AF133037; AAD24473.1; -.
EMBL; AF336825; AAK21904.1; -.
                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                    Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN-SPRAGUE DAWLEY;
Li X., Lynn B., Nagy J
"RHAMM cDNA from rat b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
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                  KOKIKHVVKLK
                                                                                                                                                                                                                                                                                                                                                                                                                        KQKIKHVVKLK
KQKI KHVVKLK
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                                                                   Similarity
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                                                                                                          836 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   713 AA;
                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                     PRELIMINARY;
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                                                                                                                                    2002) to the EMBL/GenBank/DDBJ databases AAH21427.1; -.
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 771
                                                                                                                                                                                                                       Chordata;
Rodentia;
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Rodentia;
                                                                                                          96670 MW;
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Last annotation updat
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                                                     Score 54; DB
Pred. No. 0.1
0; Mismatches
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Pred.
                                                                                                                                                                                                                     Craniata; Vert
Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Rat
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                                                                                                          B9BDDA22EBECA652 CRC64;
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thi; Muridae; Murinae; Mus
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                                                                                DB 11;
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                                                                               Length 836;
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                                                                                                                                                                                                                                                                 (RHAMM).
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Last sequence update)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local
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Q8S8K9;
01-JUN-2002
01-JUN-2002
01-JUN-2002
      STRAIN=CV.
Town C.D.,
Submitted (
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Pfam; PF00069; pkinase; 1.

PRINTS; PR00019; LEURICHPFT.

ProDom; PD000001; Euk pkinase; 1.

SMART; SM00370; LRR; 5.

SMART; SM00220; S_TKC; 1.

SMART; SM002219; TYKC; 1.

SMART; SM002219; TYCK; 1.

PROSITE; PS50011; PROTEIN KINASE DOM; 1.

ATP-binding; Kinase; Transferase.

SEQUENCE 680 AA; 74751 MW; D1079C3CF316D36D CRC64;
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Tabata S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IFR002290; Ser_thr_pkinase.
InterPro; IPR001245; Tyr_pkinase.
Pfam; PF00560. Ten
                                                                                                                                                                                                                                                                                                             STRAIN=CV. COLUMBIA;
Rounsley S.D., Lin X.,
Shen M., Ronning C.M.,
Submitted (NOV-1998) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Arabidopsis thaliana (Mouse-ear cress).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosid
eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Expressed protein. AT2G06005.
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Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Streptophyta;
                                                                                                                                                                                                                                                         SEQUENCE
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DNA Res. 7:31-63(2000)
                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                               Submitted
                                                                                                                                                                                               Lin
                                                                                                                                                                                                                         STRAIN=CV.
                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=3702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR000719; Euk_pkinase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AB018109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        features of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=20181125;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=COLUMBIA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=3702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Structural
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KQKVRHLVKL 621
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FROM N.A.
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                                                                                                                                                                                                                                                         FROM N.A.
. COLUMBIA;
, Kaul S.;
(FEB-2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IPR001611;
IPR003592;
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2 (TrEMBLrel.:
2 (TrEMBLrel.:
                                                                                                                                                                                                                            COLUMBIA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (TrEMBLrel. 21, L
ke protein kinase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRELIMINARY;
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Y., Kaneko T., K
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                                                                                                                                                                                                                                                                                                                                                 Kaul S., Shea T.P., Fuji
Fraser C.M., Somerville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21,
                                                                                                                                                                                                                                                                                                                    the
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      the
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Last sequence update)
Last annotation updat
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      EMBL/GenBank/DDBJ
                                                                                                                                                               EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                    EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 42; DB
Pred. No. 19;
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                                                                                                                                                                                                                                                                                                                                                 Fujii C.Y.,
ille C.R., V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10;
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                                                                                                                                                                                                                                                                                                                                                 Venter J.C.;
                                                                                                                                                                                                                                                                                                                                                                           Mason T.M.,
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RESULT
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ID 199
AC P9
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Q9AVX8
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Best Local S
Matches 6
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Q9AVX8

Q9AVX8

Q9AVX8

O1-JUN-2001 (TrEMBLrel. 17, Created)
O1-JUN-2001 (TrEMBLrel. 17, Last sequence up
O1-DEC-2001 (TrEMBLrel. 19, Last annotation
Hypothetical 104.0 kDa protein.
Guillardia theta (Cryptomonas phi).
Eukaryota; Cryptophyta; Cryptomonadaceae; Gu
NCBI_TaxID=55529;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-1997
01-MAY-1997
01-DEC-2001
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Eukaryota; Viridiplantae; Streptophyta; Embry,
Spermatophyta; Magnoliophyta; eudicotyledons;
Asteridae; eussterids I; Lamiales; Oleaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; AC005970; AAM15099.1; -. SEQUENCE 355 AA; 40585 MW;
MEDLINE=21223349; PubMed=11323671; Douglas S., Zauner S., Fraunholz M., Beatc Wu X., Reith M., Cavalier-Smith T., Maier "The highly reduced genome of an enslaved Nature 410:1091-1096(2001).
                                                                                                                                                                                         MEDLINE=20087226; PubMed=10618395; Zauner S., Fraunholz M., Wastl J., Pe Cavalier-Smith T., Maier U., Douglas "Chloroplast protein and centrosomal telomeres in an unusually compact euk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rosati C., Cadic A., Duron M., Renou J.P., Simoneau P.;
"Molecular cloning and expression analysis of dihydroflavonol reductase gene in flower organs of Forsythia x intermedia.";
Plant Mol. Biol. 35:303-311(1997)
EMBL; Y09127; CAA70345.1;
SEQUENCE 371 AA; 41856 MW; 4FAF14F5F4A8B366 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=CV. SPRING GLORY; TISSUE=P
MEDLINE=98009967; PubMed=9349254;
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                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
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                                                                                                                                                                        nucleomorph
                                                                                                         SEQUENCE FROM N.A.
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108 QKLKHVVRL 110
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7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
6; Conserv
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                                                                                                                                                      Acad.
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                                                                                                                                                      Sci.
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                                                                                                                                                   U.S.A.
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                                                                                                                                                                                              compact eukaryotic
                                                                                                                                                                                                                                                                                                                                                                                                                              Created)
Last sequence update)
Last annotation update)
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Pred. No. 58;
4; Mismatches
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Pred. No.
                                                                                                                                                      97:200-205 (2000)
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                                                                                                                                                                                                                                                    Penny S.L.,
                    Beaton M.,
Maier U.;
laved algal
                                                                                                                                                                                                                  genes,
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37;
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                        nucleus.";
                                                                Penny S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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RESULT 11
Q9VBU6
ID 9VBU6
AC Q9VBU
AC Q9VBU
DT 01-MA
DT 01-MA
DT 01-JU
DE CG11E
GN CG11E
GN CG1E
CN CO
CN CN CN
C
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C Q9VBU6;

TO 1-MAY-2000 (TrEMBLrel. 13, Created)

DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)

DT 01-JUN-2000 (TrEMBLrel. 21, Last annotation update

CG11857 protein (RE24638p).

GN CG11857 protein (RE24638p).

GN CG11857 protein (RE24638p).

CG Pteryota; Metazoa; Arthropoda; Tracheata; Hexap

OC Eukaryota; Neoptera; Endopterygota; Diptera; Bra

Thvdroidea; Drosophilidae; Drosophila.
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01-JUN-2001 (TrEMBLrel. 17, La
01-JUN-2001 (TrEMBLrel. 17, La
Hypothetical 16.9 kDa protein.
T30N20_260.
MEDLINE=20196006; PubMed=10731132; Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D. Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R. George R.A., Lewis S.E., Richards S., Ashburner M., Henderson Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
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Submitted (JUL-2000) to the EM
EMBL; AL365234; CAB96854.1; -
                                                                                                                                                                            SEQUENCE FROM N.A. STRAIN-BERKELEY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hypothetical sequence 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pfam; PF02519; Auxin_inducIble; 1.
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Bevan M., Peters S.A., V
Bancroft I., Mewes H.W.,
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae;

eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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01-OCT-2000
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Hypothetical protein.
SEQUENCE 861 AA; 104022 |
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ilarity 66.7%;
Conservative
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(TrEMBLrel. 13, Last sequence update)
(TrEMBLrel. 21, Last annotation update)
tein (RE24638p).
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., Rudd S., Lemcke K., Mayer K.F.X.;
o the EMBL/GenBank/DDBJ databases.
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a; Brachycera; Muscomorpha;
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Best Local
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                                                                                                        Q99QZ8 PRELIMINARY; PRT; 624 AA.
Q99QZ8; O1-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                 Staphylococcus aureus Bacteria; Firmicutes;
                                                      Staphylococcus aureus
                                                                       Hypothetical protein SAV2640 OR SA2433.
Staphylococcus
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EMBL; AE003751; AAF5
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AAM11216.1;
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23721 MW;
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                 (strain N315).
Bacillus/Clostridium
                                                      (strain Mu50 /
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Pred. No. 50;
4; Mismatches
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                                                    ATCC 700699), and
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                 group; Bacillales;
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EMBL; AP003366; BABS8802.1; -.

EMBL; AP003366; BABS8802.1; -.

EMBL; AP003366; BABS8802.1; -.

InterPro; IPR00150; BglG antitermin.

InterPro; IPR00150; PTS EIIA 2.

Pfam; PF00874; BglG antitermin; 1.

Pfam; PF00879; BglG antitermin; 1.

Pfam; PF00359; PTS EIIA 2; 1.

PROSITE; PS00372; PTS EIIA 2; 1.
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Q9AVW8;
01-JUN-2001
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Kuroda M., Ohta T., Uchiyama I., Baba T., Yuzawa H., Kobayashi I.,
Cui L., Oguchi A., Aoki K.-I., Nagai Y., Lian J.-Q., Ito T.,
Kanamori M., Matsumaru H., Maruyama A., Murakami H., Hosoyama A.,
Mizutani-Ui Y., Takahashi N.K., Sawano T., Inoue R.-I., Kaito C.,
Sekimizu K., Hirakawa H., Kuhara S., Goto S., Yabuzaki J.,
Kanehisa M., Yamashita A., Oshima K., Furuya K., Yoshino C., Shiba
Hattori M., Ogasawara N., Hayashi H., Hiramatsu K.;
"Whole genome sequencing of meticillin-resistant Staphylococcus
                                                                                                                                                        Pfam;
                                                                                                                                                              Douglas S., Zauner S., Fraunholz M., Beaton M., Penny S., I Wu X., Reith M., Cavalier-Smith T., Maier U.; "The highly reduced genome of an enslaved algal nucleus."; Nature 410:1091-1096(2001).

EMBL; AJ010592; CAC27103.1; -.
InterPro; IPR000722; RNA pol A.
InterPro; IPR0002879; RNA pol A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=20087226; PubMed=10618395; Penny S.L., Zauner S., Fraunholz M., Wastl J., Penny S.L., Cavaller-Smith T., Maier U., Douglas S.; "Chloroplast protein and centrosomal genes, a telomeres in an unusually compact eukaryotic general compact."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Guillardia theta (Cryptomonas phi).
Eukaryota; Cryptophyta; Cryptomonadaceae;
NCBI_TaxID=55529;
                                                                                                   Pfam; PF01854
DNA-directed
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01-JUN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleomorph.
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                                                                        PF00623; RNA pol A; 1.
PF01854; RNA pol A2; 1.
irected RNA polymerase; 1
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Pred. No. 1.4e+02;
2; Mismatches 2
                                                                        Nucleotidyltransferase; Transferase MW; FA556B4C96F0BBFD CRC64;
  Score
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01-MAR-2002
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01-OCT-2000
01-OCT-2000
01-DEC-2001
                                                                                                       Shimizu T. Shiba T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arabīdopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosid
eurosids II; Brassicales; Brassicaceae; Arabidopsis.
NCBI_TaxID=3702;
                                                                                                                                                                               Clostridium perfringens.
Bacteria; Firmicutes; Bacillus/Clostridium group;
Clostridiales; Clostridiaceae; Clostridium.
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                                        Hypothetical SEQUENCE 10
                                                                                                                         STRAIN=13 / TYPE A;
PubMed=11792842;
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Submitted (MAR-2000) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bevan M., Peters S.A., van Staveren M., Dirkse W., Stiekema Bancroft I., Mewes H.W., Rudd S., Lemcke K., Mayer K.F.X.; Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.
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                                                                                  flesh-eater.";
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                                                           Natl. Acad. Sci. U.S.A. AP003194; BAB82088.1; -
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Similarity 7; Conserv
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                                                                                                       Ogasawara N.,
                                                                                           Ohtani K., Hirakawa H., Ohshima K.,
Ngasawara N., Hattori M., Kuhara S., 
Jenome sequence of Clostridium perfri
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2 (TrEMBLrel. 20, 1
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31 protein CPE2382
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102 AA; 11428
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Pred. No.
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       '(cgn2_6/ptodata/2/pubpaa/US06_NEM_PUB.pep:*
'(cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
'(cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
'(cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
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Copyright (c) 1993 - 2003 Compugen Ltd
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/cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
     747
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    10 US-09-815-242-11507
9 US-09-974-143A-2
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9 US-09-815-242-10790
9 US-09-870-759-82
9 US-10-105-694-2
10 US-09-747-521-2
10 US-09-747-521-2
10 US-09-930-871-10
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Sequence 35, Appl
Sequence 11507, Appl
Sequence 11507, Appli
Sequence 2, Appli
Sequence 7, Appli
Sequence 82, Appli
Sequence 2, Appli
Sequence 10, Appli
Sequence 10, Appli
Sequence 10, Appli
Sequence 10, Appli
Sequence 18, Appli
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57.4	57.4	57.4	57.4	57.4	57.4	57.4		57.4		57.4	57.4	57.4	58.3	58.3	59.3	59.3	59.3	59.3	59.3	59.3	61.1	61.1	61.1	61.1	61.1
368	363	363	338	335	295	258	237	186	179	179	132	67	359	125	400	281	281	213	131	39	2009	1998	1973	1962	1453
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US-09-925-299-888	US-09-943-722-130	US-10-003-035-37	US-10-076-785-10	US-10-076-785-12	US-09-510-332-143	US-09-815-242-12207	US-09-815-242-5666	US-10-003-035-45	US-09-872-523-59	US-10-003-035-44	US-10-003-035-43	US-09-925-299-1310	US-09-799-777-45	US-09-925-301-933	US-09-815-242-5324	US-09-853-701-17	US-09-895-913A-282	US-09-881-752A-370	US-09-738-626-4317	US-09-864-761-46628	US-09-930-871-12	US-09-930-871-2	US-09-930-871-14	US-09-930-871-4	US-09-930-871-16
Sequence 888, App	Sequence 130, App		Sequence 10, Appl	Sequence 12, Appl	Sequence 143, App	Sequence 12207, A	Sequence 5666, Ap	Sequence 45, Appl	Sequence 59, Appl	Sequence 44, Appl	Sequence 43, Appl	Sequence 1310, Ap	Sequence 45, Appl	Sequence 933, App	Sequence 5324, Ap	Sequence 17, Appl	Sequence 282, App	Sequence 370, App	Sequence 4317, Ap	Sequence 46628, A	Sequence 12, Appl	Sequence 2, Appli	Sequence 14, Appl	Sequence 4, Appli	Sequence 16, Appl

ALIGNMENTS

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; FEATURE:
; NAME/KEY: BINDING
; LOCATION: (1)..(12)
; OTHER INFORMATION: A peptide mimicking Domain I (amino acids 401-411)
; OTHER INFORMATION: of the hyaluronic acid binding region of RHAMM
US-08-996-470-1
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                                                                                                                                                                                                                                                                                                                                                             EARLIER APPLICATION NUMBER: US 07/675,908
EARLIER FILING DATE: 1991-07-03
EARLIER APPLICATION NUMBER: US 07/838,674
EARLIER FILING DATE: 1992-02-21
EARLIER FILING DATE: 1992-02-21
EARLIER FILING DATE: 1992-02-25
EARLIER APPLICATION NUMBER: US 08/125,398
EARLIER APPLICATION NUMBER: US 08/125,398
EARLIER FILING DATE: 1993-09-23
EARLIER APPLICATION NUMBER: US 08/285,764
EARLIER APPLICATION NUMBER: US 07/952,095
EARLIER FILING DATE: 1994-08-03
EARLIER FILING DATE: 1992-09-28
NUMBER OF SEO ID NOS: 2
SOFTWARE: PATENTIN Ver: 2.0
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                        Query Match
Best Local Similarity
  Matches
                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/08/996,470A
CURRENT FILING DATE: 1997-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                            LENGTH: 1
TYPE: PRT
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  Conservative
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Score 54; DB 8; Pred. No. 0.001; Mismatches (
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FILE REFERENCE: 109272.147

CURRENT APPLICATION NUMBER: US/09/801,368

CURRENT FILING DATE: 2001-03-07

PRIOR APPLICATION NUMBER: US 09/487,558

PRIOR APPLICATION NUMBER: US 60/160,587

PRIOR FILING DATE: 1999-10-20

NUMBER OF SEQ ID NOS: 440

SOFTWARE: Patentin version 3.0

SEQ ID NO 348

LENGTH: 785

TYPE: PRT
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Best Local Similarity
6; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 2
US-09-801-368-348
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                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 34027, Application US/09864761 Patent No. US20020048763A1
                                                                                                                                                                                                                                                                     APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Aeomica-X-1
                                                                                                                                                                                                                                                                                                                               APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K
APPLICANT: Chen, Wensheng
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                                                                        PRIOR APPLICATION NUMBER: GB 24263.6 PRIOR FILING DATE: 2000-10-04
                                                                                                              PRIOR APPLICATION NUMBER: US 09/632,366 PRIOR FILING DATE: 2000-08-03
                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/207,456 PRIOR FILING DATE: 2000-05-26
                                                                                                                                                                                           CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Summers, Eric
TITLE OF INVENTION: Methods for improving Secondary Metabolite Production in Fungi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Busby, Robert APPLICANT: Cali, Brian
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                                  APPLICATION NUMBER: US 60/236,359 FILING DATE: 2000-09-27
                APPLICATION NUMBER: PCT/US01/00666
                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION:
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Silva, Jeff
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2001-01-30
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OTHER INFORMATION: MAP TO ACOOSOGO.2

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 3.2

OTHER INFORMATION: EXPRESSED IN BOULT LIVER, SIGNAL = 3.4

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 2.8

OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 3.3

OTHER INFORMATION: EXPRESSED IN LIVER, SIGNAL = 3.4

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 3.3

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 3.3

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 3.3

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 3.4

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 3.4

OTHER INFORMATION: EXPRESSED IN HELAD, SIGNAL = 3.4

OTHER INFORMATION: SWISSPROT HIT: P21439, EVALUE 1.10e-01
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US-09-840-787-35
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                                                                                                                                                                                                                                                                                                          Sequence 35, Application US/09840787
Patent No. US20020058264A1
GENERAL INFORMATION:
APPLICANT: Lal, Preeti
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SOFTWARE: Annomax Sequence Listing Engine vers.
SEQ ID NO 34027
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Best Local
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PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
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PRIOR FILING DATE: 2000-05-30
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es 6; Conserv
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00661
FILING DATE: 2001-01-30
                                                                                            Guegler, Karl J.
COTIEY, Neil C.
TITLE OF INVENTION: HUMAN REGULATORY MOLECULES
NUMBER OF SEQUENCES: 98
                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/US01/00670
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APPLICATION NUMBER: PCT,
FILING DATE: 2001-01-30
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APPLICATION NUMBER:
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                   ADDRESSEE: Incyte Pharmaceuticals, STREET: 3174 Porter Drive
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                      Yue, Henry
                                                                                                                                                                                                                                    Shah, Purvi
                                                                                                                                                                                                                                                         Hillman, Jennifer
Bandman, Olga
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NUMBER: PCT/US01/00668
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Pred. No. 6
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US-09-799-777-73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 73, Application US/09799777 Patent No. US20020091244A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        163 KRQIKHMIKYR 173
          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/799,777
FILING DATE: 06-Mar-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                             Shah, Purvi
TITLE OF INVENTION: HUMAN SIGNAL PEPTIDE-CONTAINING PROTEINS
NUMBER OF SEQUENCES: 154
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: PF-0356 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/518,865
FILING DATE: <Cnknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/840,787
FILING DATE: 23-Apr-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Lal, Preeti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
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CLONE: 1720770
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                                                                                                                                                                                                         STATE: CALIFORNIA
                                                                                                                                                                                                                                               STREET: 3174 PORTER DRIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
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CLASSIFICATION: <Unknown>
                                                                                                                                                                                         COUNTRY: USA
                                                                                                                                                                                                                             CITY: PALO ALTO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                         94304
                                                                                                                                                                                                                                                                                                                                                                                                   Hillman, Jennifer L.
Corley, Neil C.
Guegler, Karl J.
                                                                                                                                                                                                                                                                                                                                                              Baugh, Mariah
Sather, Susan
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45.5%;
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Pred. No.
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APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification of Essential Genes in
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR APPLICATION NUMBER: 60/242,578
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-815-242-11507; Sequence 11507, Application; Patent No. US20020061569A1
                                                           ; ORGANISM: Helicobacter pylori US-09-815-242-11507
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                                                                                                                                     SOFTWARE: FastSEQ
SEQ ID NO 11507
Query Match
Best Local Similarity
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                                                                                                                                                         NUMBER OF SEQ ID NOS: 14110 SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Haselbeck, Robert
                                                                                                                                                                                                PRIOR FILING DATE: 2001-02-16
                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/257,931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 73:
                                                                                                                  LENGTH: 481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          204 KRQIKHMIKYR 214
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nes 5; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 KQKIKHVVKLK 11
                                                                                                                                                                                                                   FILING DATE: 2000-12-22 APPLICATION NUMBER: 60/269,308
                                                                                                                                                                                                                                                                        FILING DATE: 2000-10-23
APPLICATION NUMBER: 60/253,625
FILING DATE: 2000-11-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/002,485 FILING DATE: <Unknown> ATTORREY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 237 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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CLONE: 3115936
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (650) 855-05
TELEFAX: (650) 845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: BILLINGS, LUCY J. REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: PF-0459 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wall, Daniel
Trawick, John D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ohlsen, Kari L.
Zyskind, Judith W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Carr, Grant J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yamamoto, Robert T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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66.7%;
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  Score 34;
Pred. No.
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Pred. No. 51;
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  DB 10;
1.1e+02;
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RESULT 9 US-09-815-242-10790 Sequence 107%, Application US/09815242
                                                                                    В
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US-09-431-226-7
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                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 7
                                                                                                                                                 Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: VERNOS, Isabelle
APPLICANT: KARSENTI Eric
APPLICANT: WITTMANN, Torster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2, Application US/09974143A Publication No. US20030022326A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 7, Application US/09431226 Patent No. US20020115599A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/431,226
CURRENT FILING DATE: 1999-11-01
PRIOR APPLICATION NUMBER: EP 99106710.9
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: KARSENTI, Eric
APPLICANT: WITTMANN, Torsten
TITLE OF INVENTION: TPX2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: JP 11/103356
PRIOR FILING DATE: 1999-04-09
NUMBER OF SEO ID NOS: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/974,143A CURRENT FILING DATE: 2002-04-19 PRIOR APPLICATION NUMBER: PCT/JP00/02281 PRIOR FILING DATE: 2000-04-07
                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1999-
                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Nezu, Jun-ichi
APPLICANT: OSE, ASUKA
TITLE OF INVENTION: NOVEL FETAL GENES
FILE REFERENCE: 06501-089001
                                                                                                                                                                                                                                               LENGTH: 747
TYPE: PRT
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TYPE: PRT
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                                                                                  131 EQKEKHHVKMK 141
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Local Similarity 63.6%;
es 7; Conservaring
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Pred. No. 1.7e+02;
"'~matches 2;
                                                                                                                                               Score 34; DB 10; Length 747; Pred. No. 1.7e+02; Mismatches 2; Indels
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US-09-870-759-82
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                                                                Matches
                                                                                               Query Match
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Best Local
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TYPE: PRT
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    144 KEKIQHVFK 152
                                1 KQKIKHVVK 9
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TITLE OF INVENTION: IDentification of Essential Genes in TITLE OF INVENTION: Prokaryotes

FILE REFERENCE: ELITRA.011A

CURRENT APPLICATION NUMBER: US/99/815,242

CURRENT APPLICATION NUMBER: 00/1-03-21

PRIOR APPLICATION NUMBER: 60/210,78

PRIOR FILING DATE: 2000-03-21

PRIOR PRIOR APPLICATION NUMBER: 60/207,727

PRIOR APPLICATION NUMBER: 60/207,727

PRIOR APPLICATION NUMBER: 60/207,727

PRIOR APPLICATION NUMBER: 60/207,727

PRIOR FILING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: 60/207,83

PRIOR APPLICATION NUMBER: 60/253,625

PRIOR APPLICATION NUMBER: 60/253,625

PRIOR FILING DATE: 2000-10-23

PRIOR APPLICATION NUMBER: 60/257,931

PRIOR APPLICATION NUMBER: 60/257,931

PRIOR APPLICATION NUMBER: 60/259,308

PRIOR FILING DATE: 2000-12-22

PRIOR PRIOR PRICATION NUMBER: 60/269,308

PRIOR FILING DATE: 2000-12-22

PRIOR PRICATION NUMBER: 60/269,308

PRIOR FILING DATE: 2000-12-22

PRIOR PRICATION NUMBER: 60/269,308

PRIOR FILING DATE: 2001-02-16

SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                  ; SOFTWARE: PatentIn version 3.1

; SEQ ID NO 82

; LENGTH: 330

; TYPE: PRT

; ORGANISM: Staphylococcus aureus

US-09-870-759-82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Enterococcus faecalis US-09-815-242-10790
                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: TERMAN, David S
Best Local Similarity Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 82, Application US/09870759 Patent No. US20020177551A1
                                                                                                                                                                                                                                                     TITLE OF INVENTION: COMPOSITIONS AND METHOUS FILLE REFERENCE: 870759

CURRENT APPLICATION NUMBER: US/09/870,759

CURRENT FILING DATE: 2002-01-14

PRIOR APPLICATION NUMBER: US 60/208,128

PRIOR FILING DATE: 2000-05-30

NUMBER OF SEQ ID NOS: 166
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Zyskind, Judith W.
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Trawick, John D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yamamoto, Robert T.
Xu, H. Howard
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  Conservative
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                        61.1%;
66.7%;
                        Score 33; DB 9;
Pred. No. 1.1e+02;
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Pred. No. 34;
Mismatches
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                                             Length 330;
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RESULT 13
US-09-747-521-2
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TYPB: PRT
; ORGANISM: Bacillus anthracis
US-10-105-694-2
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US-10-105-694-2
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US-10-105-695-2
GENERAL INFORMATION:
APPLICANT: Galloway, Darrel
APPLICANT: Meteczun, Alfred
APPLICANT: Meteczun, Served
TITLE OF INVENTION: Methods for Protection Against Lethal Infection with Bacillus
FILE REFERENCE: 22727/04079
CURRENT APPLICANTON NUMBER: US/09/747,521
CURRENT FILING DATE: 2000-12-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Galloway, Darrel R.
APPLICANT: Mateczun, Alfred J.
APPLICANT: Mateczun, Alfred J.
TITLE OF INVENTION: Methods for Protection Against Lethal Infection with Bacillus FILE REFERENCE: 22727/04115
CURRENT APPLICATION NUMBER: US/10/105,695
CURRENT FILING DATE: 2002-03-25
CURRENT FILING DATE: 2002-03-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/10105694 Publication No. US20030003109A1 GENERAL INFORMATION:
                                                                                                                                        Sequence 2, Application US/09747521 Patent No. US20020051791A1
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LENGTH: 809
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/105,694
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: US 09/747,521
PRIOR FILING DATE: 2000-12-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Galloway, Darrel R.
APPLICANT: Mateczun, Alfred J.
TITLE OF INVENTION: Methods for Protection Against Lethal Infection with Bacillus
FILE REFERENCE: 22727/04116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 09/747,521
PRIOR FILING DATE: 2000-12-21
NUMBER OF SEQ ID NOS: 8
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Similarity 45.5%;
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Pred. No. 2
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Pred. No. 2.7e+02;
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2.7e+02;
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Query Match
Best Local Similarity
Trahes 5; Conserve
; NAME/KEY: VARIANT
LOCATION: (1)...(1381)
; OTHER INFORMATION: Xaa =
US-09-930-871-8
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; Sequence 2, Application US
; Patent No. US20020142002A1
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                                                                                                                                           SEQ ID NO 8
                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/225,989
PRIOR FILING DATE: 2000-08-16
NUMBER OF SEQ ID NOS: 20
SOFTWARF.
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APPLICANT: Mateczun, Alfred J.
TITLE OF INVENTION: Methods for Protection Against Lethal Infection with Bacillus FILE REFERENCE: 22727/04114
CURRENT APPLICATION NUMBER: US/10/106,014
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: US 09/747,521
PRIOR APPLICATION NUMBER: US 09/747,521
PRIOR FILING DATE: 2000-12-21
NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                               FILE REFERENCE: LEX-0216-USA
CURRENT APPLICATION NUMBER: US/09/930,871
CURRENT FILING DATE: 2001-08-14
                                                                                                                                                                                                                                                                                                                     APPLICANT: Turner, C. Alexander Jr. APPLICANT: Mathur, Daniel APPLICANT: Mathur, Brian
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SOFTWARE: PatentIn version 3.0
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                                                                                                                                                          SOFTWARE:
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TYPE: PRT
ORGANISM: Bacillus anthracis
                                                                   ORGANISM: homo sapiens FEATURE:
                                                                                                      LENGTH: 1381
TYPE: PRT
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FastSEQ for Windows Version 4.0
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Pred. No. 2.7e+02;
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Patches 6: Conservative 1: Mismatches 1: Indels 0: Caps 0:

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Search completed: February 8. 2003, 10:52:09

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54
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Copyright (c) 1993 - 2003 Compugen Ltd.
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/cgn2_6/ptodata/1/paa/US080_COMB.pep:*
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US-08-318-892B-3

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Sequence 3, Appli
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Sequence 4, Appli
Sequence 30, Appl
Sequence 30, Appl
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Sequence 47, Appl	47,	4, 7	80,	`		77,	18,	14,	16,	e 16,	48,	48,		673	e 88	2, Ap	79,		e 42	2, Appli	80	33, 1	78,	e 83,	73,	463,	74, 1	Sequence 463, App	76	7 5	02,	0 C	51,	e 51,	e 1, App	Sequence 9, Appli	0

ALIGNMENTS

RESULT 1 US-08-318-892-3

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Sequence 3, Application US/08318892

GENERAL INFORMATION:
APPLICANT: TUTLEY, EVA A
TITLE OF INVENTION: HYALURONAN RECEPTOR (RHAMM = RECEPTOR TITLE OF INVENTION: HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING TITLE OF INVENTION: FOR HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING TITLE OF INVENTION: FOR HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING TITLE OF INVENTION: FOR HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING TITLE OF INVENTION: PEPTIDES

NUMBER OF SEQUENCES: 29

CORRESPONDENCE ADDRESS:

ADDRESSE: C/O FISH & NEAVE
STREET: 1251 Avenue of the Americas
CITY: New York
STATE: N.Y.
COUNTRY: USA
ZIP: 10020

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATION SYSTEM: PC-DOS/MS-DOS
SOCTWARE: WOORDERICECT 6.1

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/318,892
FILLING DATE: 1-DEC-1994
CLASSIFICATION NUMBER: US/08/318,892
FILLING DATE: 1-DEC-1994
REGISTRATION NUMBER: 30,709
REFERENCE/DOCKET NUMBER: 30,709
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RESULT 3
US-09-169-077-1
; Sequence 1, Application US/09169077
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GENERAL INFORMATION:
APPLICANT: TUTLEY, EVA A
TITLE OF INVENTION: HYALURONAN RECEPTOR
TITLE OF INVENTION: HYALURONAN MEDIATED
                                                                                                                                                                Matches
                                                                                                                                                                                                Query Match
                                                                                                                                                              Best Local Similarity Matches 11; Conserv
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Best Local Similarity
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                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (212) 596-90 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Plerri, Margaret A
REGISTATION NUMBER: 30709
REFERENCE/DOCKET NUMBER: SIM-8
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPOTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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ADDRESSEE: c/o FISH & NEAVE
STREET: 1251 Avenue of the Americas
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TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/318,892B
FILING DATE: 1-DEC-1994
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                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
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Pred. No. 0.0071;
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RESULT 5
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; MOLECULE TYPE:
US-09-169-077-1
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                                                                                                                                                                                                                                                                   SEQ ID NO 4
LENGTH: 11
TYPE: PRT
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GENERAL INFORMATION:
APPLICANT: Turley, Eva A
TITLE OF INVENTION: Improved Delivery of Disease Modifiers
FILE REFERENCE: 205375
                                                                                                                                                 Query Match 100.0%;
Best Local Similarity 100.0%;
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Best Local Similarity
                                                                                                                                   Matches
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FILING DATE: 10-APR-1996
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 1997-05-22 NUMBER OF SEQ ID NOS: 36
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                                                                                                                                                                                                                                    FEATURE:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/169,077
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MEDIUM TYPE: Floppy disk
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NUMBER OF SEQUENCES:
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STRANDEDNESS: sir
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Pred. No. 0.0071;
Mismatches 0
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Sequence 30, Application US/09685010; GENERAL INFORMATION:
APPLICANT: Turley, Eva A.
APPLICANT: Cruz, Tony F.

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US-08-318-892-9
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US-09-978-309A-30
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            Sequence 9, Application US/08318892
GENERAL INFORMATION:
APPLICANT: Turley, Eva A
TITLE OF INVENTION: HYALURONAN RECEPTOR (RHAMM = RECEPTOR
TITLE OF INVENTION: FOR HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING
TITLE OF INVENTION: PEPTIDES
NUMBER OF SEQUENCES: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATING
TITLE OF INVENTION: CELLULAR RESPONSE TO INJURY AND OTHER PROLIFERATING CELL
TITLE OF INVENTION: DISORDERS REGULATED BY HYALADHERIN AND HYALJRONANS
FILE REFERENCE: 910130.401C1
CURRENT APPLICATION NUMBER: US/09/685,010
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 72
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 30
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LENGTH: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated TITLE OF INVENTION: Hyaladherin and Hyaluronans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Cruz, Tony
APPLICANT: Pastrak, A
APPLICANT: Turley, E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/978,309A CURRENT FILING DATE: 2002-04-04
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Peptide composition that binds a hyalauronan
                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
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CORRESPONDENCE
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Local Similarity 100.0%;
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Turley, Eva A.
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Pred. No. 0.0071;
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Pred. No. 0.0071;
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              TELEFAX: (212) 596-9090 INFORMATION FOR SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (212) 596-90 INFORMATION FOR SEQ ID NO:
                                                       REGISTRATION NUMBER: 30709
REFERENCE/DOCKET NUMBER: SI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 1-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Pierri, Margaret A
                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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NAME: Pierri, Margaret A
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 SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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STREET: 1251
CITY: New Yor
                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0
                                                                                                                                                                                                                                                                                                         CITY: New York STATE: N.Y.
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                                            TELEPHONE:
                                                                                                                                                                                                                                                                                       COUNTRY:
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                                                                                                                                                                                                PatentIn Release #1.0, Version #1.30
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EARLIER FILING DATE: 1992-02-21
EARLIER APPLICATION NUMBER: US 07/838,675
EARLIER FILING DATE: 1992-02-21
EARLIER APPLICATION NUMBER: US 08/125,398
EARLIER APPLICATION NUMBER: US 08/285,764
EARLIER APPLICATION NUMBER: US 08/285,764
EARLIER FILING DATE: 1994-08-03
EARLIER APPLICATION NUMBER: US 07/952,095
EARLIER FILING DATE: 1992-09-28
NUMBER OF SEQ ID NOS: 2
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                           APPLICANT: Turley, Eva A.
APPLICANT: Cruz, Tony F.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATING
TITLE OF INVENTION: CELLULAR RESPONSE TO INJURY AND OTHER PROLIFERATING
TITLE OF INVENTION: DISORDERS REGULATED BY HYALADHERIN AND HYALURONANS
FILE REFERENCE: 910130.401C1
CURRENT APALICATION NUMBER: US/09/685,010
CURRENT FILING DATE: 2001-05-23
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COTHER INFORMATION: A peptide mimicking Domain I (amino acids 401-411)
OTHER INFORMATION: of the hyaluronic acid binding region of RHAMM
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Pred. No. 0.0079;
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RESULT 12
US-09-978-309A-81
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SOFTWARE: Fas
SEQ ID NO 51
FENGTH: 32
                                                                                                                                                                                                                                                        Sequence 81, Application US/09978309A GENERAL INFORMATION:
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Best Local Similarity
Matches 11; Conserv
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Best Local Similarity 100.
Matches 11; Conservative
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APPLICANT: Turley, Eva A.

APPLICANT: Turley, Eva A.

APPLICANT: Turley, Eva A.

TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response to
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated by
TITLE OF INVENTION: Hyaladherin and Hyaluronans
FILE REFERENCE: 03352-010

CURRENT APPLICATION NUMBER: US/09/978,309A

CURRENT APPLICATION NUMBER: US 09/685,010

PRIOR APPLICATION NUMBER: US 09/685,010

PRIOR APPLICATION NUMBER: US 09/541,522

PRIOR APPLICATION NUMBER: US 09/541,522

PRIOR FILING DATE: 2000-04-03

PRIOR FILING DATE: 2000-04-03
                                                                                                    APPLICANT: Cruz, Tony
APPLICANT: Pastrak, Aleksandra
APPLICANT: Turley, Eva A.
APPLICANT: Turley, Eva A.
TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response to
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated by
TITLE OF INVENTION: Hyaladherin and Hyaluronans
FILE REFERENCE: 033352-010
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PRIOR APPLICATION NUMBER: US 09/685,010 PRIOR FILING DATE: 2000-10-05 PRIOR APPLICATION NUMBER: US 09/541,522
                                           CURRENT APPLICATION NUMBER: US/09/978,309A CURRENT FILING DATE: 2002-04-04 PRIOR APPLICATION NUMBER: US 09/685,010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/127,457
PRIOR FILING DATE: 1999-04-01
NUMBER OF SEQ ID NOS: 84
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TYPE: PRT
ORGANISM: Unknown
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ORGANISM: Unknown
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Pred. No. 0.025;
Mismatches
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Pred. No. 0.025;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; ORGANISM: Mus musculus US-09-978-309A-82
                                                                                                                                                                                                                                                                                                      US-09-657-276-1174
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                                                                                                                                                                                                                                                                Sequence 1174, Application US/09657276 GENERAL INFORMATION:
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Best Local Similarity
Matches 11; Conserv
                                                                         TITLE OF INVENTION: ITTLE OF INVENTION: ITTLE OF INVENTION: CEPTLE REFERENCE: 2110
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SOFTWARE: FASTSEQ for Windows Version 4
SEQ ID NO 82
LENGTH: 32
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LENGTH: 32
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 09/685,010
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: US 09/541,522
PRIOR FILING DATE: 2000-04-03
PRIOR APPLICATION NUMBER: US 60/127,457
PRIOR FILING DATE: 1999-04-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Turley, Eva A.
TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response to
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated by
TITLE OF INVENTION: Hyaladherin and Hyaluronans
FILE REFERENCE: 033352-010
CURRENT APPLICATION NUMBER: US/09/978,309A
CURRENT FILING DATE: 2002-04-04
CURRENT APPLICATION NUMBER: US/09/657,276
CURRENT FILING DATE: 2000-09-07
PRIOR APPLICATION NUMBER: 60/134,406
PRIOR FILING DATE: 1999-05-17
                                                                                                                                                                                     APPLICANT:
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APPLICANT: Pastrak, A
APPLICANT: Turley, E
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TYPE: PRT
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PRIOR FILING DATE: 1999-04-01
NUMBER OF SEQ ID NOS: 84
SOFTWARE: FastSEQ for Windows Version 4
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Local Similarity 100.0%;
nes 11; Conservative (
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                                                                                                                                                Milner, Peter
Holmes, Darren
Thibaudeau, Karen
                                                                                                                                                                                                           Bridon, Dominique
Ezrin, Alan
                                                                                                                                                                                                                                                 Conjuchem, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                              PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM PEPTIDASE ACTIVITY THROUGH CONJUGATION TO BLOOD
                                                                                          COMPONENTS
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Pred. No. 0.025;
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Pred. No. 0.025;
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; LENGTH: 221
; TYPE: PRT
; ORGANISM: MUS N
US-09-978-309A-75
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Search completed: February Job time: 147.3 secs
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                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 09/685,010
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: US 09/541,522
PRIOR FILING DATE: 2000-04-03
PRIOR APPLICATION NUMBER: US 60/127,457
PRIOR FILING DATE: 1999-04-01
NUMBER OF SEQ ID NOS: 84
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 75, Application US/09978309A GENERAL INFORMATION: APPLICANT: Cruz, Tony
                                                                                                                                                                                                                                                                                       SEQ ID NO 75
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SEQ ID NO 1174
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Best Local Similarity
                                                                                                                                          Best Local Similarity
Matches 11; Conserva
                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Pastrak, Aleksandra
APPLICANT: Turley, Eva A.
TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response to
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated by
TITLE OF INVENTION: Hyaladherin and Hyaluronans
FILE REFERENCE: 03352-010
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PRIOR FILING DATE: 1999-09-10
PRIOR APPLICATION NUMBER: 60/1
PRIOR FILING DATE: 1999-10-18
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                     173 KQKIKHVVKLK 183
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                                                                                      1 KQKIKHVVKLK 11
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                                                                                                                                          100.0%; ilarity 100.0%; Conservative 0;
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Pred. No. 0.
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OM protein - protein search, using sw model
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2003, 10:41:35; Search time 16.5 Seconds (without alignments)
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Sequence: Title: Perfect score: US-09-685-010-30 54 1 KQKIKHVVKLK 11

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched

405691 seqs, 81078759 residues

Total number of hits satisfying chosen parameters:

Maximum DB seq length: 0
DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Pending_Patents_AA_New:*

1: /cgn2_6/ptodata/2/paa/U
2: /cgn2_6/ptodata/2/paa/U
3: /cgn2_6/ptodata/2/paa/U
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/cgn2_6/ptodata/2/paa/USO1_NEW_COMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Result No.	Score	Query Match	Length	DB	ID	Description
1	37	68.5	272	ۍ ا	US-09-134-000C-6100	Sequence 6100, Ap
2	37	68.5	272	ű	-09-134-000C-61	6100,
ω	37	68.5	624	S	-09-950-084-	Œ
4	36	66.7	234	6	-10-092-411A-319	3195,
5	36	66.7	284	5	-09-134-000C-	e 5495,
6	36	66.7	284	5	-549	5495,
7	36	66.7	1388	۳	PCT-US02-21338-164	164,
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9	35	64.8	188	Ç	-084-480	4804,
10	35	64.8	708	G	24-676-9635	96355
11	35	64.8	708	σ	53	96355,
12	35	64.8	715	տ	-09-724-676-963	e 96353,
13	35	64.8	715	5	635	96354,
14	35	64.8	715	ഗ	4-676A-963	Sequence 96353, A
15	35	64.8	715	5	US-09-724-676A-96354	96354,
16	35	64.8	797	5	-724-676-9635	96352,
17	35	64.8	797	5	US-09-724-676A-96352	96352,
18	. 35	64.8	804	σ	US-09-724-676-96351	96351,
19	35	64.8	804	S	US-09-724-676A-96351	
20	35	64.8	1097	σ	-77	2203,
21	34	63.0	30	6	US-10-203-138A-11061	e 11061,
22	34	63.0	405	6	-092-	5259,
23	34	63.0	433	σ	231	
24	34	63.0	433	σ	US-09-724-676A-52315	Sequence 52315, A
25	34	63.0	518	6	-10-092-411A-474	4744, A
26	34	63.0	588	σ	US-09-724-676-52313	52313,

45	44	43	42	41	40	39	38	37	36	ა ე	34	33	32	31	30	29	28	27
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US-10-304-095-8	US-10-245-882-210	US-09-724-676A-52314	US-09-724-676-52314	PCT-US02-29560-210	US-10-228-063-36	US-10-228-063-26	PCT-US02-27129-36	PCT-US02-27129-26	US-10-228-063-27	PCT-US02-27129-27	US-10-228-063-5	US-10-191-715-2	PCT-US02-27129-5	US-09-724-676A-71216	US-09-724-676A-71215	US-09-724-676-71216	US-09-724-676-71215	US-09-724-676A-52313
Sequence 8, Appli	Sequence 210, App	Sequence 52314, A	Sequence 52314, A	Sequence 210, App			Sequence 36, Appl	Sequence 26, Appl	Sequence 27, Appl	Sequence 27, Appl	Sequence 5, Appli	Sequence 2, Appli	Sequence 5, Appli	Sequence 71216, A	Sequence 71215, A	Sequence 71216, A	Sequence 71215, A	Sequence 52313, A

ALIGNMENTS

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Sequence 6100, Application US/09134000C

Sequence 6100, Application US/09134000C

GENERAL INFORMATION:

APPLICANT: Lynn Doucette-Stamm et al
APPLICANT: Lynn Do
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US-09-134-000C-6100
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US-09-134-000C-6100
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                                                                                                                               Sequence 6100, Application US/09134000C

GENERAL INFORMATION:

APPLICANT: Lynn Doucette-Stamm et al

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO

TITLE OF INVENTION: ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 032796-032

CURRENT APPLICATION NUMBER: US/09/134,000C

CURRENT FILING DATE: 1998-08-13

PRIOR APPLICATION NUMBER: US 60/055,778

PRIOR APPLICATION NUMBER: US 60/055,778

PRIOR SEQ ID NOS: 6812

SOFTWARE: Patentin version 3.1

SEQ ID NO 6100

LENGTH: 272
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                                                                                                 LENGTH: 2
TYPE: PRT
                                        ORGANISM: Enterococcus faecalis
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US-10-092-411A-3195; Sequence 3195, Application US/10092411A
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US-09-950-084-6869
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Staphylococcus aureus US-09-950-084-6869
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NUMBER OF SEQ ID NOS: 7451
SEQ ID NO 6869
LENGTH: 624
TYPE: PRT
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APPLICANT:
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                                                          APPLICANT: Lynn Doucette-Stamm et al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: 032796-101 CURRENT APPLICATION NUMBER: US/10/092,411A CURRENT FILING DATE: 2002-03-07 PRIOR APPLICATION NUMBER: US 09/134,001 PRIOR EPILING DATE: 1998-08-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Roberta S. Hare
APPLICANT: Karen J. Shaw
TITLE OF INVENTION: Staphylococcus aureus Related Compositions and Methods
FILE REFERENCE: 1034/1C963US2
CURRENT APPLICATION NUMBER: US/09/950,084
CURRENT FILING DATE: 2001-09-10
                      PRIOR A&PLICATION NUMBER: US 60/064,964 PRIOR R/IL*NG DATE: 1997-11-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 09/036,338 PRIOR FILING DATE: 1998-03-06
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PRIOR APPLICATION NUMBER: US 09/353,718
PRIOR FILING DATE: 1999-07-14
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APPLICATION NUMBER: US 09/266,541
FILING DATE: 1999-03-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 09/266,555
FILING DATE: 1999-03-11
APPLICATION NUMBER: US 09/266,542
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FILING DATE: 1999-03-11
APPLICATION NUMBER: US 09/266,556
FILING DATE: 1999-03-11
APPLICATION NUMBER: US 60/055,779
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US-09-134-000C-5495

Sequence 5495, Application US/09134000C

GENERAL INFORMATION:

APPLICANT: Lynn Doucette-Stamm et al

APPLICANT: Lynn Doucette-Stamm et al

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACI

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACI

TITLE REFERENCE: 032796-032

CURRENT APPLICATION NUMBER: US/09/134,000C

CURRENT FILING DATE: 1998-08-13

PRIOR APPLICATION NUMBER: US 60/055,778

PRIOR APPLICATION NUMBER: US 60/055,778

PRIOR FILING DATE: 1997-08-15

NUMBER OF SEQ ID NOS: 6812

SOFTWARE: Pattentin version 3.1

SEQ ID NO 5495

LENGTH: 284

TYPE: PRT
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Best Local Similarity
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                                                                                                                                                       ; ORGANISM: Enterococcus faecalis US-09-134-000C-5495
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                                                                                                                                                                                                                                                                                                                             Sequence 5495, Application US/09134000C
GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al
APPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 032796-032
CURRENT APPLICATION NUMBER: US/09/134,000C
CURRENT FILING DATE: 1998-08-13
CURRENT FILING DATE: 1998-08-13
                                                                                                                                                                                                            CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/055,778
PRIOR FILING DATE: 1997-08-15
NUMBER OF SEQ ID NOS: 6812
SOFTWARE: Patentin version 3.1
SEQ ID NO 5495
LENGTH: 284
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NUMBER OF SEQ ID NOS: 5676
SEQ ID NO 3195
LENGTH: 234
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Matches 6; Conserv
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TYPE: PRT
                                                                                                                                                                                                TYPE: PRT
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244 KENIEHVVRVK 254
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KENIEHVVRVK 254
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                                                                            Conservative
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77.8%;
                                                                                           Score 36;
Pred. No.
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Pred. No.
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                                                                                         DB
26;
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21;
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                                                                                                               Length 284;
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TITLE OF INVENTION: Cancer
FILE REFERENCE: 018501-002330US
CURRENT APPLICATION NUMBER: US/10/188,832
CURRENT FILING DATE: 2002-11-22
PRIOR APPLICATION NUMBER: US 60/302,814
PRIOR FILING DATE: 2001-07-03
PRIOR APPLICATION NUMBER: US 60/310,099
PRIOR FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: US 60/343,705
PRIOR FILING DATE: 2001-11-13
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; ORGANISM: Homo sapiens
US-10-188-832-164
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                                                                                                   SOFTWARE: Pa
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Best Local
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LENGTH: 1388
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PRIOR FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: US 60/343,705
PRIOR FILING DATE: 2001-11-08
PRIOR APPLICATION NUMBER: US 60/350,666
PRIOR APPLICATION NUMBER: US 60/350,666
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                                                                                                                                         NUMBER OF SEQ ID NOS: 207
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TITLE OF INVENTION: Methods of Diagnosis of Bladder Cancer, Compositions
TITLE OF INVENTION: and Methods of Screening for Modulations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Mack, David H. APPLICANT: Aziz, Natasha APPLICANT: Eos Biotechno
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CURRENT FILING DATE: 2002-12-04
PRIOR APPLICATION NUMBER: US 60/302,814
PRIOR FILING DATE: 2001-07-03
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TITLE OF INVENTION: Methods of Diagnosis of Bladder Cancer, Compositions
TITLE OF INVENTION: and Methods of Screening for Modulators of Bladder
TITLE OF INVENTION: Cancer
FILE REFERENCE: 018501-002330PC
                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/372,246 PRIOR FILING DATE: 2002-04-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
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NUMBER OF SEQ ID NOS: 207
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                                                    LENGTH: 1388
TYPE: PRT
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                                                                                                                                 PatentIn Ver.
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Pred. No. 1.5e+02;
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RESULT 10
US-09-724-676-96355
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Matches
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LENGTH: 188
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                                                         SEQ ID NO 96355
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Best Local
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                                                                      APPLICANT: Compugen LTD TITLE OF INVENTION: Variants of alternative splicing FILE REFERENCE: 129181.4 Compugen CURRENT APPLICATION NUMBER: US/09/724,676 CURRENT FILING DATE: 2000-11-28 NUMBER OF SEQ ID NOS: 97222 SOFTWARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Remaining Prior Applicatio
NUMBER OF SEQ ID NOS: 7451
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PRIOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 1999-10-14
PRIOR APPLICATION NUMBER: US 09/353,718
PRIOR FILING DATE: 1999-07-14
PRIOR APPLICATION NUMBER: US 09/266,557
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 09/036,338 PRIOR FILING DATE: 1998-03-06
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TYPE: PRT
ORGANISM: Homo sapiens
                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
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FILLING DATE: 1999-03-11
APPLICATION NUMBER: US 09/266,556
FILING DATE: 1999-03-11
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Roberta S. Hare
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66.7%;
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70.0%;
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Pred. No. 1.5e+02;
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RESULT 13
US-09-724-676-96354
US-09-724-676-96354, Application US/09724676
Sequence 96354, Application US/09724676
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 12914: 4 Compugen
CURRENT: APPLICATION NUMBER: US/09/724,676
CURRENT: TILING DATE: 2000-11-28
NUMBER OF *FO ID NOS: 97222
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TITLE OF INVENTION: Variants of alternative spirite REFERENCE: 129181.4 Compugen

CURRENT APPLICATION NUMBER: US/09/724,676A

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 9722

SOFTWARE: Patentin version 3.2

SEQ ID NO 96355

LENGTH: 708

TYPE: PAT

ORGANISM: Homo sapiens

US-09-724-676a-96355
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US-09-724-676-96353
Sequence 96353, Application US/09724676
GENERAL INFORMATION:
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US-09-724-676A-96355
; Sequence 96355, Application US/09724676A
; GENERAL INFORMATION:
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Best Local Similarity
These 7; Conserve
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US-09-724-676-96353
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TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patentin version 3.2
SEQ ID NO 96353
LENGTH: 715
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Best Local Similarity
Matches 7; Conserv
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Matches 7; Conserv
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127 KIPHVVRLK 135
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127 KIPHVVRLK 135
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77.8%;
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77.8%;
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Pred. No. 1.1e+02;
Prematches 1;
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Pred. No. 1.1e+02;
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Pred. No. 1.1e+02;
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SOFTWARE: PatentIn version 3.2

SEQ ID NO 96353

LENGTH: 715

TYPE: PRT

ORGANISM: Homo sapiens
US-09-724-676A-96353
                                                                                                            Search completed: February Job time: 18.5 secs
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                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patentin version 3.2
SOFTWARE: Patentin version 3.2
LEUGTH: 715
TYPE: PRT
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APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
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Best Local :
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Pred. No. 1.1e+02;
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US-08-477-831C-31
US-08-477-831C-3195
US-09-134-001C-3195
US-09-732-191-2
US-09-733-750C-35
US-09-734-613-35
US-09-134-001C-5259
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US-08-402-217A-2

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US-08-477-831C-38
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Sequence 5144, Ap	Sequence 96, Appl	Sequence 17, Appl	Sequence 9, Appli	Sequence 72, Appl	Sequence 53, Appl	Sequence 1, Appli	Sequence 1, Appli	Sequence 7, Appli	Sequence 83, Appl	Sequence 2, Appli	Sequence 2, Appli	Sequence 2, Appli	•	Sequence 10, Appl	Sequence 10, Appl	Sequence 8, Appli	Sequence 8, Appli

ALIGNMENTS

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; DESCRIPTION: US-08-477-831C-38
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                                                                              Query Match
Best Local
                                                               Matches
                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-596-9000
TELEPAX: 212-596-9090
INFORMATION FOR SEQ ID NO: 38:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 10020-1104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICANT:
APPLICANT:
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 1251 AVE
CITY: NEW YORK
STATE: NEW YORK
COUNTRY: U.S.A.
ZIP: 10020-1104
                                                                                                                                                                                                                                                                                                                     NAME: PIERRI, MARGARET A. REGISTRATION NUMBER: 30,7 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                             Local
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                1 KQKIKHYVKLK 11
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KQKIKHVVKLK 11
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5. 6429291
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11; Conserv
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SHUWEN, ZHANG
ENTWISTLE, JOYCELYN
VENTION: HYALURONAN RECEPTOR PROTEIN
                                                            100.0%; ilarity 100.0%; Conservative (
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                                                            Score 54; DB 4; Pred. No. 0.0039; Mismatches 0
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                                                                                            Length 32;
                                                               Indels
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US-08-477-831C-41

Sequence 4 Patent No.

1, Application US/08477831C 6429291

GENERAL INFORMATION:

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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 183 amino acids
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COMPUTER READABLE FORM:
REDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/NG-DOS
                                                                                                     TITLE OF INVENTION: HYALURONAN RECEPTOR EXPRESSED IN TITLE OF INVENTION: UMBILICAL VEIN ENDOTHELIAL CELLS NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS: ADDRESS: INCYTE PHARMACEUTICALS, INC.
                                                                                                                                                                                          APPLICANT: Hawkins, Phillip R. APPLICANT: Wilde, Craig G. APPLICANT: Seilhamer, Jeffrey J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Rel. #1.0, ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,831C
FILING DATE: 07-JUN 1995
CLASSIFICATION: 514
COMPUTER KSADABLE FORM:
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ZIP: 10020-110
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APPLICANT: STUDEN, ZHANG
APPLICANT: ENTWERN, ZHANG
APPLICANT: ENTWERN, ZHANG
TITLE OF INVENTION: HYALURONAN RECEPTOR PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: (CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                             166 KQKIKHVVKLK 176
                                   COUNTRY:
                                                         STAZE:
                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: PIERRI, MARGARET A.
REFERENCE/DOCKET NUMBER: SIM-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 1251 AVIOLETY: NEW YORK STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 100.0%; es 11; Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: sir
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                                                                                                                                                                                                                                                                                                                                                                                                               1 KQKIKHVVKLK 11
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                  94304
                                                                  Palo Alto
                                                     CA
                                                                                      3174 Porter Drive
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                                    USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4; Length 183;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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; SEQUENCE CHARACTERISTICS:

; LENGTH: 351 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

US-08-402-217A-2
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Best Local Similarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                      APPLICATION NUMBER: US/08/700,178
FILING DATE: August 20, 1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/402,217
FILING DATE: March 10, 1995
ATTORNEY/ACENT INFORMATION:
                                 TELEFAX: 415-845-4166 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                              NAME: Billings, Lucy J. PREGISTRATION NUMBER: 36,749 REFERENCE/DOCKET NUMBER: PETELECOMMUNICATION INFORMATION: TELEPHONE: 415-855-9555
                    SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Hawkins, Phillip R.
APPLICANT: Wilde, Craig G.
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: HYALURONAN RECEPTOR EXPRESSED IN
TITLE OF INVENTION: UMBILICAL VEIN ENDOTHELIAL CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION.
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  318 KQKIKHVVKLK 328
                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: INCLLE Drive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: 10-MAR
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351 amino acids
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SYSTEM: PC-DOS/MS-DOS
Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                        August 20, 1996
ON: 435
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TOPOLOGY:
US-08-995-654-2
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RESULT 6
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                                                                                                                                          Query Match
Best Local :
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GENERAL INFORMATION:
                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/995,654
FILING DATE: December 22, 1997
                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 351 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: PF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/700,178
FILING DATE: August 20, 1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Dr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/402,217 FILING DATE: March 10, 1995 ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Hawkins, Phillip R. APPLICANT: Wilde, Craig APPLICANT: Seilhamer, Jeffrey
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: HYALURONAN RECEPTOR EXPRESSED IN HUMAN TITLE OF INVENTION: UMBILICAL VEIN ENDOTHELIAL CELS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
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CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                          TELEFAX: 650-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: UZIP: 94304
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                                                                                    1 KOKIKHVVKLK 11
                                                                                                                                       Similarity
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                                                                                                                         Conservative
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                                                                                                                                                                                                                linear
                                                                                                                                       100.0%; Score 54; DB 3; Length 351; 100.0%; Pred. No. 0.043;
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                                                                                                                         Mismatches
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; LIBRARY: mouse
; CLONE: GI 53979
US-08-402-217A-3
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                                                                                                                                                                                                                                                      Sequence 3, Application US/08700178 Patent No. 5783669 Patent No. 5783669 5700912
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100
Best Local Similarity 100
Matches 11; Conservative
                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Hawkins, Phillip R.
APPLICANT: Wilde, Craig G.
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: HYALURONAN RECEPTOR EXPRESSED IN
TITLE OF INVENTION: UMBILICAL VEIN ENDOTHELIAL CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DDS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Hawkins, Phillip R. APPLICANT: Wilde, Craig G. APPLICANT: Seilhamer, Jeffrey J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION: NAME: Luther, Barbara J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                        CORRESPONDENCE ADDRESS:
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                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                      402 KOKIKHVVKLK 412
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                                             STREET: 3174 Por
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Luther, Barbara J
REGISTRATION NUMBER: 339
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 3174 Por
CITY: Palo Alto
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TOPOLOGY: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: INCYTE PHARMACE STREET: 3174 Porter Drive
COUNTRY: U
ZIP: 94304
                                    STATE: CA
                                                                                       ADDRESSEE:
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                                                                    3174 Porter Drive
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                                                                                         INCYTE PHARMACEUTICALS, INC
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Pred. No. 0.058;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 477;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3, Application US/08995654
Patent No. 6025138
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 477 anino acids
TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-055
APPLICATION NUMBER: 08/700,178
FILING DATE: August 20, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/402,217
FILING DATE: March 10, 1995
                                                                                                                                                                    COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/402,217
FILING DATE: March 10, 1995
ATTORNEY/ACENT INFORMATION:
NAME: Billings, Lucy J.
                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: HYALURONÂN RECEPTOR EXPRESSED IN HUMAN TITLE OF INVENTION: UMBILICAL VEIN ENDOTHELIAL CELS NUMBER OF SEQUENCES: 3
                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Wilde, Craig APPLICANT: Seilhamer, Jeffrey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IMMEDIATE SOURCE:
LIBRARY: mouse
CLONE: GI 53979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 402 KOKIKHVVKLK 412
                                                                                                                                                                                                                                                                                                                           STREET: 3174 POI
CITY: Palo Alto
STATE: CA
                                                                                                                                   APPLICATION NUMBER: US/08/995,654 FILING DATE: December 22, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: August 20, 1996
CLASSIFICATION: 435
TOR ADDITION: 435
                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 100 nes 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Incyte Pharmaceuticals, Inc
STREET: 3174 Porter Dr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
TOPOLOGY: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
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Pred. No.
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Query Match
Best Local Similarity
Thes 11; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: TURLEY, EVA A.
APPLICANT: SHUMEN, ZHANG
APPLICANT: ENTWLSTLE, JOYCELYN
TITLE OF INVENTION: HYALURONAN RECEPTOR PROTEIN
                                                                                                                                                                                                                         TELEPHONE: 212-596-900
TELEFAX: 212-596-9090
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 477 amino acids
                                                                                                     MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/477,831C
FILING DATE: 07-70U1-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: PIERRI, MARGARET A.
REGISTRATION NUMBER: 30,709
REFERENCE/DOCKET NUMBER: SIM-10
                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Rel. #1.0, ASCII
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 650-855-0555
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NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-
                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS
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CITY: NEW YORK
STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Match 100.0%; Score 54; DB 3; Local Similarity 100.0%; Pred. No. 0.058;
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                                                                                                                                                                                        LENGTH:
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                                                                                                                                                                      amino acid
                                                                                                                                                                                        606 amino acids
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                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                         linear
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linear
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                                                                                                  protein
/desc = "RHAMM I protein"
                                                                                                                                                   single
                                   100.0%;
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                     0;
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                   Score 54; DB 4; Length 606; Pred. No. 0.074; o; Mismatches 0; Indels
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                   Gaps
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531 KQKIKHVVKLK 541

RESULT 10 US-08-477-831C-11

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RESULT 11
US-08-477-831C-40
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                                                                                                                                                                                                                  Sequence 40, Application US/08477831C Patent No. 6429291 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS_MS-DOS
SOFTWARE: Patentin Rel. #1.0, ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,831C
FILING DATE: 07-JUN-1995
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
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INFORMATION FOR SEQ ID NO: 11:
                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: FISH & NEAVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 631 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: PIERRI, MARGARET A.
REGISTRATION UNMBER: 30,7
REFERENCE/DOCKET NUMBER:
                                                                                                                                             APPLICANT: TURLEY, EVA A.
APPLICANT: SHUMEN, ZHANG
APPLICANT: ENTWISTLE, JOYCELYN
TITLE OF INVENTION: HYALURONAN RECEPTOR PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: SHUWEN, ZHANG
APPLICANT: ENTWISTLE, JOYCELYN
TITLE OF INVENTION: HYALURONAN RECEPTOR PROTEIN
                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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STATE: NEW YORK
COUNTRY: U.S.A.
ZIP: 10020-1104
                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
STATE: NEW YORK COUNTRY: U.S.A. ZIP: 10020-1104
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                                                       CITY: NEW YORK
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                                                                      1251 AVENUE OF THE AMERICAS
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1251 AVENUE OF THE AMERICAS
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US-08-477-831C-39
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Best Local S
Matches 10
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                TELEFAX: 212-596-9090 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                          COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC pDS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Rel. #1.0, ASCII CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/477,831C
                                                                                                                                                           CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: PIERRI, MARGARET A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: TURLEY, EVA A APPLICANT: SHUWEN, ZHANG APPLICANT: ENTWISTLE, JOURNALL OF INVENTION: HYALL
                                                                                               REGISTRATION NUMBER: 30,709
REFERNCE/DOCKET NUMBER: SI
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: SI
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 07-JUN-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: DIEBBY MARCABET A
                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                               SOFTWARE: Patentin Rel. #1.0, ASCII CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                           STREET: 1251 AVENCITY: NEW YORK
STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    166 KQKIRHVVKLK 176
                                                                TELEPHONE: 212-596-9090
                                                                                                                                                                                                  APPLICATION NUMBER: US/OFILING DATE: 07-JUN-1995 CLASSIFICATION: 514
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A. ZIP: 10020-1104
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STRANDEDNESS:
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                             LENGTH:
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10; Conserv
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6429291
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                           32 amino acids
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VENTION: HYALURONAN RECEPTOR PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TURLEY, EVA A.
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N: 514
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US-09-572-191-2
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; DESCRIPTION: /desc = protein fragment
US-08-477-831C-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Staphylococcus epidermidis US-09-134-001C-3195
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US-09-134-001C-3195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 3195, Application US/09134001C
Patent No. 6380370
GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCCCCUS
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCCCCUS
TITLE REFERENCE: GTC-007
CURRENT APPLICATION NUMBER: US/09/134,001C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 3195
LENGTH: 234
TYPE: DET
                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/572,191
CURRENT FILING DATE: 2000-05-17
NUMBER OF SEQ ID NOS: 6
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 1388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 81.0
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Matches 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2, Application US/09572191 Patent No. 6355466
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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APPLICANT: Sakowicz, Roman
APPLICANT: Wood, Kenneth
TITLE OF INVENTION: No. 6355466el motor proteins and methods for
TITLE OF INVENTION: their use
                                                                                                                                                                                                                                                                                      FILE REFERENCE:
                                                                                                                                                                     TYPE: PRT
210 KPKMKHVVK 218
                                 2 OKİKHVVKLK 11
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77.88;
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81.8%;
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                                                                              Score 36; DB 4;
Pred. No. 2.1e+02;
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Search completed: February Job time: 13.4667 secs

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Best Local Similarity The
Conservat
                                                                                                                                                    ; ORGANISM: Human US-09-723-262-2
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                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 1388
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/09723262 Patent No. 6379912 GENERAL INFORMATION:
                                                                                                                                                                                                                                                            APPLICANT: Beraud, Christophe
APPLICANT: Sakowicz, Roman
APPLICANT: Wood, Kenneth
TITLE OF INVENTION: No. 6379912el motor proteins and methods for
TITLE OF INVENTION: their use
TITLE REFERENCE: 1017
CURRENT APPLICATION NUMBER: US/09/723,262
CURRENT APPLICATION NUMBER: US/09/723,262
CURRENT FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: US 09/572,191
PRIOR APPLICATION NUMBER: US 09/572,191
NUMBER OF SEQ ID NOS: 6
1351 QKIQYVVRLK 1360
                  2 QKIKHVVKLK 11
|||::||:||
                                                                          Conservative
                                                                                              66.7%;
70.0%;
                                                                       Score 36; DB 4; Le
Pred. No. 2.1e+02;
3; Mismatches 0;
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                                                                            Indels
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GenCore version 5.1.3 Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: February 8, 2003, 10:39:10 ; Search time 12 Seconds (without alignments)
80:112 Million cell updato

cell updates/sec

Sequence: Title: Perfect score: 1 KLKSQLVKRK 10 US-09-685-010-31

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Total number of hits satisfying chosen parameters:

283224 seqs, 96134422 residues

283224

Minimum DB Maximum DB seq length: 0 seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : **4** ω PIR_73:*
1: pir1:*
2: pir2:*
3: pir3:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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ALIGNMENTS

TACA298 hyaluronan receptor - mouse (7.Species: Mus musculus (house mouse) (7.Species: Mus musculus (house mouse) (7.Species: 16-Nov-1995 #sequence_revision 08-Feb-1996 #text_change 01-Dec-2000 (7.Accession: JC4298; A42925; Ā41923; S21.586 R.EINTWISTLE, J.; Zhang, S.; Yang, B.; Wong, C.; Li, Q.; Hall, C.L.; A, J.; Mowat, M. A; Fittle: Characterization of the murine gene encoding the hyaluronan receptor RHAMM. A; Reference number: JC4298; MUID:96011639; PMID:7590272 A; Molecule type: mRNA A; Asesidues: 1-631 ERUT> A; Cross-references: EMBL:X64550 A; Experimental source: 3T3 fibroblast R; Hardwick, C. J. Cell Biol. 118, 753, 1992 A; Reference number: A42925; MUID:92348516; PMID:1639856 A; Contents: erratum A; Accession: A42925 A; Molecule type: mRNA A; Molecule type: mRNA A; Molecule type: mRNA
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A;Residues: 155-376,'S',378-504,'E',506-631 <HAR>
A;Cross-references: GB:X64550

A;Note: authors translated the codon AGT for residue 377 as Thr and CTA for residue 507 a R;Hardwick, C.; Hoare, K.; Owens, R.; Hohn, H.P.; Hook, M.; Moore, D.; Cripps, V.; Auster J. Cell Biol. 117, 1343-1350, 1992
A;Title: Molecular cloning of a novel hyaluronan receptor that mediates tumor cell motiliar, Reference number: A41923; MUID:92299690; PMID:1376732

A;Status: preliminary A;Molecule type: mRNA A;Residues: 155-376,'S',378-504,'E',506-507,'L',508-630 <HA2>

A;Cross-references: GB:X64550 A;Note: this sequence has been corrected in reference A42925 C;Comment: This protein regulates cell motility and transformation, and focal adhesion di

A;Gene: rhamm
A;Gene: rhamm
A;Gene: rhamm
A;Introns: 20/3; 54/2; 79/2; 137/3; 187/3; 259/2; 382/2; 431/2; 482/2; 515/3; 574/3; 625/C;Superfamily: hypaluronan receptor
C;Keywords: glycoprotein; receptor
F;260-382/Region: 21 residue repeats
F;516-594/Region: 21 residue repeats
F;516-574/Region: hyaluronan binding #status predicted
F;575-625/Region: hyaluronan binding #status predicted
F;515-625/Region: hyaluronan binding #status predicted
F;116,283,304,325,346,367,398,438,619/Binding site: carbohydrate (Asn) (covalent) #status

Query Match Best Local Similarity 90.0 Warches 9; Conservative 93.5%; Score 43; DB 2; 90.0%; Pred. No. 1; Mismatches 0, Length 631; Indels

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A;Reference number: A86625; MUID:21235186; PMID:11337471
A;Accession: A86657
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C;Species: Lactococcus lactis subsp. lactis
C;Date: 23-Mar-2001 #sequence_revision 23-Mar-2001 #text_change 11-Jan-2002
C;Accession: A86657
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Pathway: gluconeogenesis
C;Superfamily: Bacillus subtilis fructose-bisphosphatase; phosphoesterase core homology
C;Keywords: carbohydrate metabolism; gluconeogenesis; metalloprotein; phosphoric monoes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Bolotin, A.; Wincker, P.; Mauger, S.; Jaillon, O.; Malarme, K.; Weissenbach, Genome Res. 11, 731-753, 2001
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aldehyde reductase (BC 1.1.1.21) - rabbit (fragment) C; Species: Oryctolagus cuniculus (domestic rabbit) C; Date: 22-Jun-1990 #sequence_revision 22-Jun-1990 # C; Acceseion: A34406 R; Garcia-Perez, A.; Martin, B.; Murphy, H.R.; Uchida J. Biolagasha. 264, 16815-16821, 1989
                                                                                                                                                                                                                                                                                                                                                                                                            A;Molecule type: DNA
A;Residues: 1-336 <LET>
A;Residues: 1-336 <LET>
A;Cross-references: EMBL:U97194; PIDN:AAB52452.1; GSPDB:GN00019; CESP:C37A2.
A;Experimental source: strain Bristol N2; clone C37A2
C;Genetics: C250
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A; Residues: 1-640 <STO>
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A; Accession: T30161
A; Status: prelimina:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Le, T.T.; Kemp, K.; Scheet, P. submitted to the EMBL Data Library, April A;Description: The sequence of C. elegans A;Reference number: Z20746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein C37A2.8 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #to
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Introns: 42/3; 77/3; 198/1; 264/1; 295/3
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                        H.; Cowley Jr.,
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RESULT 5
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                                                                                                                                                                                                                                                                                                                                                                        A;Molecule type: protein
A;Roelecule type: protein
A;Roelecule type: protein
A;Roelecule type: part of this sequence was confirmed by protein sequencing
R;Graham, C.B.; Szpirer, C.; Levan, G.; Carper, D.
Gene 107, 259-267, 1991
A;Title: Characterization of the aldose reductase-encoding gene famil
A;Reference number: I53649; MUID:92084118; PMID:1748296
A;Accession: I53649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Carper, D.; Nishimura, C.; Shinohara, T.; Dietzchold, B.; FEBS Lett. 220, 209-213, 1987
A;Title: Aldose reductase and rho-crystallin belong to the A;Reference number: S00079; MUID:87276556; PMID:3111886
A;Accession: S00079.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Carper, D.A.; Wistow, G.; Nishimura, C.; Graham, C.; Watanabe, K.; Fujii, Y.; Hayashi, Exp. Eye Res. 49, 377-388, 1989
A;Title: A superfamily of NADPH-dependent reductases in eukaryotes and prokaryotes.
A;Reference number: A60603, MUID:90005742; PMID:2507340
A;Accession: A60603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Species: Rattus norvegicus (Norway rat)
C;Date: 20-Apr-2000 #sequence revision 20-Ap
C;Accession: A60603; S00079; E60603; IS3649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: GB:J05048; NID:g164756; C;Superfamily: aldehyde reductase C;Keywords: oxidoreductase
                                                                                                                         C;Keywords: -
F;263/Active
                                                                                                                                                                                                                                            A;Cross-references: GB:M60322; NID:g202851; PIDN:AAA40721.1; PID:g202852 C;Comment: Aldose reductase catalyzes reduction of a variety of sugars to sugar alcohols C;Comment: This enzyme is active in the eye lens, where an accumulation of sugar alcohol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Molecule type: mRNA
A;Residues: 33-316 <CA2>
A;Cross-references: EMBL:X05884
A;Note: 45-Ser and 54-Trp were also
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A; Residues: 1-302 <GAR>
                                                                                                                                C;Superfamily: aldehyde reductase
C;Keywords: eye lens; NADP; oxidoreductase
F;263/Active site: Lys #status predicted
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KLKSQLVKRK 10
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Rigui, T.; Tanimoto, T.; Kokai, Y.; Nishimura, C. Eur. J. Biochem. 227, 448-453, 1995
A;Title: Presence of a closely related subgroup in the aldo-ketoreductase family of the A;Reference number: I49484; MUID:95154325; PMID:7851421
A;Accession: I49484
A;Status: preliminary; translated from GB/EMBL/DDBJ
hypothetical protein BBH39 - Lyme disease spirochete plasmid H/lp28-3 C;Species: Borrella burgdorferi (Lyme disease spirochete) C;Date: 13-Feb-1998 #sequence_revision 13-Feb-1998 #text_change 08-Oct-1999 C;Accession: E70239
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A;Residues: 94-174,'N',176-177,'D',179-181,'DRKPVVYSCYRPSWSPSQRRCIKHTSDKLIKGFLHLFKNLFV'
A;Cross-references: EMBL:D89260; NID:g1749727; PIDN:BAA13921.1; PID:g1749728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA Res. 4, 363-369, 1997

A;Title: Identification of open reading frames in Schizosaccharomyces A;Reference number: Z17323; MUID:98162722; PMID:9501991

A;Accession: T43185
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R;Wood, V.; Rajandream, M.A.; Barrell, B.G.; Bothe, G.; Pohl, T. submitted to the BMBL Data Library, December 1998
A;Reference number: Z21955
A;Accession: T40898
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A;Experimental source: strain 972h-; cosmid c1259
R;Yoshioka, S.; Kato, K.; Nakai, K.; Okayama, H.; Nojima, H.
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                                                                                                                      RESULT 8
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ily: pyruvate dehydrogenase complex protein X; lipoyl/biotin-binding homology
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A;Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser, A;Title: The complete genome sequence of the gastric pathogen Helicobacter pylori.
A;Reference number: A64520; MUID:97394467; PMID:9252185
A;Accession: B64527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       penicillin-binding protein dacf precursor (dacF) RP389 - Rickettsia prowazekii C;Species: Rickettsia prowazekii C;Date: 21-Nov-1998 #sequence_revision 21-Nov-1998 #text_change 03-Nov-2000 C;Accession: D71696 R;Andersson, S.G.E.; Zomorodipour, A.; Andersson, J.O.; Sicheritz-Ponten, T.; Alsmark Nature 396, 133-140, 1998 A;Title: The genome sequence of Rickettsia prowazekii and the origin of mitochondria. A;Reference number: A71630; MUID:99039499; PMID:9823893 A;Accession: D71696
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A;Title: Genomic sequence of a Lyme disease spirochaete,
A;Ritle: Genomic sequence of a Lyme disease spirochaete,
A;Reference number: A70100; MUID:98065943; PMID:9403685
A;Accession: E70239
A;Status: preliminary; nucleic acid sequence not shown;
                                                                                                                                                                       R;Tomb, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; Fleischmann, R.D. Peterson, S.; Loftus, B.; Richardson, D.; Dodson, R.; Khalak, H.G.; Glodek, A.; McKenney son, J.D.; Kelley, J.M.; Cotton, M.D.; Weidman, J.M.; Fujii, C.; Bowman, C.; Watthey, L. Nature 388, 539-547, 1997

A;Authors: Wallin, B.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser, C.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: GB:AJ235271; GB:AJ235269; NID:g3868717; PIDN:CAA14846.1; PID:g386094; A;Experimental source: strain Madrid E
C;Genetics:
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                       A; Molecule type: DNA
A; Residues: 1-362 < TOM>
                                                                                                                                                                                                                                                                                                   M-protein - Helicobacter pylori (strain 26695)
C;Species: Helicobacter pylori
C;Species: Helicobacter pylori
C;Date: 30-Jan-1998 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998
C;Accession: B64527
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A; Residues: 1-306 < AND>
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A; Residues: 1-34 < KLE>
                                                                    A;Status: preliminary; nucleic
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GB:AE000511; TIGR:HP0058
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Pred. No.
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Pred. No. 3
                                                                         sequence not
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3.7;
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                                                                              shown; translation
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Score

34;

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362;

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RESULT 13
                                                  retrovirus-related pol polyprotein pseudogene - human
N;Contains: endonuclease (EC 3.1.-.-); RNA-directed DNA polymerase (EC 2.7.
C;Species: Homo sapiens (man)
C;Date: 28-Dec-1987 #sequence_revision 04-Jan-1996 #text_change 14-May-1999
C;Accession: D24483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;anonymous, Genoscope
submitted to the EMBL Data Library, July 1999
s,Description: Pyrococcus abyssi genome sequence:
                   R;Ono, K.; Yasunaga, T.; Mi
J. Virol. 60, 589-598, 1986
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A; Accession: D75112
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C;Species: Pyrococcus abyssi
C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Date: 21-Jan-2000 #sequence_revision 21-Jan-2000 #text_change 21-Jul-2000 C;Accession: T28323
R;Afonso, C.L.; Tulman, E.R.; Lu, Z.; Oma, E.; Kutish, G.F.; Rock, D.L.
J. Virol. 73, 533-552, 1999
A;Title: The genome of Melanoplus sanguinipes entomopoxvirus.
A;Reference number: Z20484; MUID:99102612; PMID:9847359
A;Accession: T28323
                                                                                                                                                                                                                                                                                                                                                      A; Gene:
                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: GB:AJ248285; GB:AL096836; NID:g5458067; PIDN:CAB49613.1; A;Experimental source: strain Orsay
                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: DNA
A; Residues: 1-633 < KAW>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C; Accession:
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A; Residues: 1-522 < AFO>
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 Nucleotide sequence of
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70.0%;
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Pred. No. 56;
3; Mismatches
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2; Mismatches
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                                    Ushikubo,
endogenous retrovirus genome related to the
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 mouse
                       A;Molecule type: DNA
A;Residues: 1-2145 <POW>
A;Residues: 1-2145 <POW>
A;Cross-references: EMBL:274219; NID:g1431273; PIDN:CAA98745.1; PID:g1431274; MIPS:YDL17
A;Cross-references: SGD:S0002330; MIPS:YDL171c
                                                                                                                      A; Reference number: A; Accession: S67723
                                                                                                                                                                           R; Pohl,
                                                                                                                                                                                                         A; Molecule type: DNA
A; Residues: 1-2145 < POH>
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                                                                                                                                                                                                                                                            A; Reference number: S61010
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A; Residues: 1-874 < ONO >
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les 7; Conserv
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                   SGD:GLT1
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7; Conserv
                                                                                                                                          S67708
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C;Species: Saccharomyces cerevisiae
C;Date: 15-Feb-1996 #sequence_revision
C;Accession: S61041; S67723
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A;Residues: 1-1785 <DEL>
A;Cross-references: GB:M34947; NID:g160597; PID:g160598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Deleersnijder, W.; Hendrix, D.; Bendahman, N.; Hanegreefs, Mol. Blochem. Parasitol. 43, 231-244, 1990
A;Title: Molecular cloning and sequence analysis of the gene A;Reference number: A45546; MUID:91218805; PMID:2090945
A;Accession: A45546
submitted to the Protein Sequence
                                          A;Cross-references: EMBL:Z67750; NID:g1061256; PIDN:CAA91574.1; PID:g1061267
                                                                                                                                                                                                                                                                           glutamate synthase (NADH2) (EC 1.4.1.14) glt1 precursor N;Alternate names: protein D1448; protein YDL171c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change
C;Accession: A45546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    major merozoite surface antigen precursor - Plasmodium chabaudi C;Species: Plasmodium chabaudi chabaudi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Keywords: endonuclease; hydrolase; nucleotidyltransferase; polyprotein;
F;36-250/Domain: RNA-directed DNA polymerase <REV>
F;585-764/Domain: endonuclease <END>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Reference number: A93023; MUID:87036922; PMID:3021993 A;Accession: D24483
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77.8%;
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  Database,
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A;Map position: 4L
C;Superfamily: glutamate synthase (NADH)
C;Koywords: 3Fe-45; metalloprotein; oxidoreductase; transmembrane protein
F;1-53/Domain: propeptide #status predicted <PRO>
F;54-2145/Product: glutamate synthase #status predicted <MAT>
F;1077-1093/Domain: transmembrane #status predicted <TM1>
F;1172-1188/Domain: transmembrane #status predicted <TM2>
F;54/Active site: Cys #status predicted
F;1185,1191,1196/Binding site: 3Fe-4S cluster (Cys) (covalent) #status predicted
F;1185,1191,1196/Binding site: 3Fe-4S cluster (Cys) (covalent) #status predicted
P;1185,1191,1196/Binding site: 3Fe-4S cluster (Cys) (covalent) #status predicted
P;185,1191,1196/Binding site: 3Fe-4S cluster (Cys) (covalent) #status predicted
P;185,1191,1191,1196/Binding site: 3Fe-4S cluster (Cys) (covalent) #status predicted
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Cell 95:591-59
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"The hyaluronan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-99059494; PubMed-9845361;
Hofmann M., Assmann V., Fieber C.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EXTRACELLULAR-REGULATED KINASE (ERK) ACTIVITY.
SUBUNIT: SUBUNIT OF THE HARC COMPLEX.
SUBCELLULAR LOCATION: CELL SUFFACE AND CYTOPLASMIC.
ALTERNATIVE PRODUCTS: 2 ISOFORMS; RHAMMIY4 (SHOWN FRHAMMI; ARE PRODUCED BY ALTERNATIVE SPLICING.
TISSUE SPECIFICITY: UBIQUITOUSLY EXPRESSED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FUNCTION: INVOLVED IN CELL MOTILITY. WHEN HYALURONA HAME, THE PHOSPHORYLATION OF A NUMBER OF PROTEINS, FOCAL ADHESION KIMASE OCCURS. MAY ALSO BE INVOLVED TRANSFORMATION AND METASTASIS FORMATION, AND IN REG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MGI:104667; Hmmr.
MGI:dacid; Alternative
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  transient tyrosine kinase activity.";
. 126:575-588(1994).
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RESULT 3
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Matches 8
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Matches
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Hyaluronic acid; Repeat; Glyc
DOWAIN 420 430
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CARBOHYD 262 262
CARBOHYD 302
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CARBOHYD 483 483
SEQUENCE 498 AA; 57858 MW;
                                                                                                                                                                                                                                                                                                                                                                             HMMR RAT
P97779;
30-MAY-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HMMR OR IHABP OR RHAMM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
30-MAY-2000 (Rel. 39, Last annotation update)
Hyaluronan mediated motility receptor (Intracellular hyaluronic acid
binding protein) (Receptor for hyaluronan-mediated motility).
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                                                                                                   1 KLKSQLVKRK 10
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                                                                                                                                                                                                                                                                                                                                                                    EXTRACELLULAR-REGULATED KINASE (ERK) ACTIVITY (BY SUBUNIT: SUBUNIT OF THE HARC COMPLEX.
SUBCELLULAR LOCATION: CELL SURFACE AND CYTOPLASMIC
                                                                                                                                                                                                                                                                                                                                                           SIMILARITY)
                                                                               KLRSQLAKRK 451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KLKSQLVKRK 10
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8; Conser
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1; Mismatches
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Pred. No.
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Query Match
Best Local :
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Oxidoreductase;
INIT_MET 0
MOD_RES 1
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ACT SITE
CONFLICT
                                                                                                                                                                                                                                                                                           PROSITE;
PROSITE;
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MEDLINE=95154325; PubMed=7851421;
Gui T., Tanimoto T., Kokai Y., Nishimura
"Presence of a closely related subgroup
family of the mouse.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                  InterPro; IPR001395; Aldo/ket_r
Pfam; PF00248; aldo_ket_red; 1.
PRINTS; PR00069; ALDKETRDTASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                    PMMA-2DPAGE; P45376;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; D32250; BAA06980.1; -.
EMBL; L39795; AAA62176.1; -.
EMBL; U29152; AAA69958.1; -.
HSSP; P15121; 2ACQ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entities requires a license agreement (Su or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Iwata T., Carper D.;
Submitted (MAR-1995) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=ICR
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15-JUN-2002 (Rel. 41, Last annotation update)
Aldose reductase (EC 1.1..1.21) (AR) (Aldehyde
AKR1B1 OR ALDR1 OR ALDOR1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-FEB-1996 (Rel.
15-JUN-2002 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                              MGD; MGI:1353494; Akr1b1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Daoudal S., Berger M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM
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61
                                    щ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBUNIT: MUNUTUMN: Cytoplasmic.
SUBCELLULAR LOCATION: Cytoplasmic.
THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VARIETY OF CARBONYL-CONTAINING COMPOUNDS TO THEIR CORRESPONDING ALCOHOLS WITH A BROAD RANGE OF CATALYTIC EFFICIENCIES.

CATALYTIC ACTIVITY: Alditol + NAD(P)(+) = aldose + NAD(P)H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SIMILARITY: BELONGS TO THE ALDO/KETO REDUCTASE FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUBUNIT: MONOMER.
                KLKSQLVKRK 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SWISS-PROT entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the EW
                                                                                                                                                                                                                               PDD000288; Aldo/ket_red; 1.
; PS00062; ALDOKETO_REDUCTASE_2; 1.
; PS00063; ALDOKETO_REDUCTASE_3; 1.
; PS00798; ALDOKETO_REDUCTASE_1; 1.
; PS00798; ALDOKETO_REDUCTASE_1; 1.
ductase; NADP; Acetylation; Multigene family.
TO 0
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Biochem. 227:448-453 (1995)
                                                                         Similarity 7; Conser
                                                                                                                                                     48
45
315 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Swiss;
                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE=Kidney;
Targer M., Pailhoux E., Tournaire
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       P45376; MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE=Liver;
                                                                                                                                                       35601 MW;
                                                                                                76.1%;
70.0%;
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                                                                                            .0%;
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                                                                                              Score 35;
Pred. No.
                                                                                                                                                                        HYDROGEN-BOND DONOR (BY A -> S (IN REF. 1).
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                                                                                                                                                       607E604FF432580E CRC64;
                                                                           Mismatches
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                                                                                                                Length 315;
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                                                                           Indels
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RESULT
ALDR_RA
pfam, prio248; aldo_ket_red; I.
priNTS; prio066; ALDKETEDIASE.
proDom; pD000288; Aldo/ket_red; 1.
proSITE; p800062; ALDOKETO_REDUCTASE_3; 1.
proSITE; p800063; ALDOKETO_REDUCTASE_3; 1.
proSITE; p8000998; ALDOKETO_REDUCTASE_1; 1.
Oxidoreductase; NADP; Acetylation.
INIT_MET
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                                                                                                                                                                                                                                                                                                                                                EMBL; U13694; AAB60687.1; JOINED. EMBL; U13699; AAB60687.1; JOINED. EMBL; U13690; AAB60687.1; JOINED. EMBL; U13691; AAB60687.1; JOINED. EMBL; U13692; AAB60687.1; JOINED. EMBL; U13692; AAB60687.1; JOINED. EMBL; U13693; AAB60687.1; JOINED. EMBL; U13216; AAA50833.1; -. EMBL; U12316; AAA51157.1; -.
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MEDLINE=95024191; PubMed=7938022;
Perraris J.D., Williams C.K., Martin B.M., Burg M.B., Garcia-Perez
"Cloning, genomic organization, and osmotic response of the aldose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-1996 (Rel. 34, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation updat Aldose reductase (EC 1.1.1.21) (AR) (Aldehy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Garcia-Perez A., Martin B., Murphy H.R., Uchida S., Murer H., Cowley B.D. Jr., Handler J.S., Burg M.B.; "Molecular cloning of cDNA coding for kidney aldose reductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Ci
Mammalia; Eutheria; Lagomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-APR-1990 (Rel.
01-OCT-1996 (Rel.
15-JUN-2002 (Rel.
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Eukaryota; Metazoa; Chordata; C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AKR1B1
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SEQUENCE
                                                                                                                                                                                                                                                                                                                             PIR; A34406; A34406.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=89380313; PubMed=2506183;
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                                                                                                                                                                                                                                               SSP; P15121; ZALV.
InterPro; IPR001395; Aldo/ket_re
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4
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FUNCTION: CATALYZES THE NADPH-DEPENDENT REDUCTION OF A WIDE VARIETY OF CARBONYL-CONTAINING COMPOUNDS TO THEIR CORRESPONDING ALCOHOLS WITH A BROAD RANGE OF CATALYTIC EFFICIENCIES.

CATALYTIC ACTIVITY: Alditol + NAD(P)(+) = aldose + NAD(P)H.

SUBCELLULAR LOCATION: Cytoplasmic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SIMILARITY: BELONGS TO THE ALDO/KETO REDUCTASE FAMILY.
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35632
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                       ACETYLATION (BY SIMILARITY). HYDROGEN-BOND DONOR (BY SIMILARITY)
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64D53E6AC0853FFB CRC64;
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                                                                                                This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outset the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for comentities requires a license agreement (See http://www.isb-sib.ch/an or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                             Laeng P., Bouillon P., Taupenot L., Labourdette G.;

"Long-term induction of an aldose reductase protein by basic
fibroblast growth factor in rat astrocytes in vittro.";

Electrophoresis 16:1240-1250(1995).

-I- FUNCTION: CATALYZES THE NADPH-DEPENDENT REDUCTION OF A WIDE
VARIETY OF CARBONYI-CONTAINING COMPOUNDS TO THEIR CORRESPONDING
ALCOHOLS WITH A BROAD RANGE OF CATALYTIC EFFICIENCIES.

-I- CATALYTIC ACTIVITY: Alditol + NAD(P)(+) = aldose + NAD(P)H.
                                                 EMBL; X05884; CAA29308.1; -. EMBL; M60322; AAA40721.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=Astrocytes;
MEDLINE=96007949; PubMed=7498172;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
MEDLINE=92084118; PubMed=1748296;
Graham C.E., Szpirer C., Levan G.,
"Characterization of the aldose re
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Craft C., Kador P., Kinoshita J.H.;
"Aldose reductase and p-crystallin belong
superfamily as aldehyde reductase.";
FEBS Lett. 220:209-213(1987).
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01-NOV-1995 (Rel. 32, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Aldose reductase (EC 1.1.1.21) (AR) (Aldehyde AKR1B1 OR ALDR1.
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                                  PIR; S00079; S00079.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gene
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MEDLINE=87276556; PubMed=3111886;
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rattus norvegicus (Rat).
                  HSSP; P15121;
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                                                                                                                                                                                                                         SUBUNIT: MONOMER.

SUBCELLULAR LOCATION: Cytoplasmic.

SUBCELLULAR LOCATION: Cytoplasmic.

DISBASE: IN DIABETES AND GALACTOSEMIA, INCREASED AR ACTIVITY LEADS
TO HIGH LEVELS OF SORBITOL AND GALACTITOL, RESPECTIVELY, IN THE
CELLS OF MANY TISSUES. ACCUMULATION OF SUGAR ALCOHOLS HAS BEEN
SHOWN TO CAUSE OSMOTIC CATARACTS IN THE LENS. AR IS ALSO THOUGHT
TO PLAY A KEY ROLE IN DIABETIC COMPLICATIONS OF THREE OTHER TARGET
TISSUES, NAMELY, NERVE, KIDNEY AND RETINA.

SIMILARITY: BELONGS TO THE ALDO/KETO REDUCTASE FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      107:259-267(1991).
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IPR001395; Aldo/ket_red
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70.0%;
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Pred.
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Best Local Similarity
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P10266; Q14273;
01-MAR-1989 (Rel
                                                                                                                      Pfam; PF00075; rnaseH; 1.
Pfam; PF00078; rvt; 1.
Pfam; PF00552; integrase; 1.
Pfam; PF00565; rve; 1.
Pfam; PF02022; Integrase Zn; 1.
Pfam; PF02022; Integrase; Zn; 1.
Rya-directed DNA polymerase; Nuclease; Endonumerase; Nuclease; Nuc
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                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; M14123; AAA88033.1; ALT_SEQ. PIR; D24483; GNHUER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-MAR-1999 (Rel. 10, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Endogenous retrovirus HERV-KlO putative pol polyprotein [Includes:
Reverse transcriptase (EC 2.7.7.49); Endonuclease].
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01-MAR-1989 (Rel.
15-JUN-2002 (Rel.
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PRINTS; PR00069; ALDKETEDTASE.
ProDom; PD000288; Aldo/ket red; 1.
PROSITE; PS00062; ALDOKETO_REDUCTASE_2;
PROSITE; PS00798; ALDOKETO_REDUCTASE_3;
PROSITE; PS00798; ALDOKETO_REDUCTASE_1;
Oxidoreductase; NADP; Acetylation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the mouse mammary tumor virus genome.";
J. Virol. 60:589-598(1986).
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Ono M., Yasunaga T., Miyata T., U
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                 DOMAIN
                                                                                                                                                                                                                                                                                              InterPro; IPR000477; RVTse.
InterPro; IPR001584; Rve.
                                                                                                                                                                                                                                                                                                                                              InterPro;
                                                                                                                                                                                                                                                                                                                                                                                           InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ono M., Yasunaga T., Miyata T., Ushikubo "Nucleotide sequence of human endogenous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                   interPro;
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IPR003308; Integrase_Zn.
IPR002156; RNaseH.
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HYDROGEN-BOND DONOR (BY SIMILARITY);
572941A154BC1202 CRC64;
                                                                                       REVERSE TRANSCRIPTASE ENDONUCLEASE.
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34;
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  (See http://www.isb-sib.ch/announce/
                                                                                                                                    Endonuclease;
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REMBL; Z67750; CAA91574.1; ...

REMBL; Z74219; CAA98745.1; ...

REMBL; Z74219; CAA98745.1; ...

RINGL; Z74219; CAA98745.1; ...

RPAGM; PF00070; pyr_redox; 1.

RPAGM; PF01493; DUF14; 1.

RPAGM; PF01493; DUF14; 1.

RPAGM; PF01493; DUF14; 1.

RPAGM; PF01645; Glu synthase; 1.

RPAGM; PF01645; Glu synthase; 1.

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GIT1 YEAST STANDARD; PRT; 21
Q12680; Q12290;
01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence up
01-NOV-1997 (Rel. 35, Last annotation
30-MAY-2000 (Rel. 39, Last annotation
                                CONFLICT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                                                                   Glutamate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (NOV-1995) to the EMBL/GenBank/DDBJ databases -i- CATALYTIC ACTIVITY: 2 L-glutamate + NADP(+) = L-glut
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=S288c;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Sequence of the GLT1 gene from Saccharomyces cerevisiae domain structure of yeast glutamate synthase."; Yeast 12:1359-1366(1996).
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Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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   SEQUENCE
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IRON-SULFUR (3FE-4S)
IRON-SULFUR (3FE-4S)
                             IPS -> FLV (IN REF. L -> V (IN REF. 2).
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FMN (BY SIMILARITY).
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(EC 1.4.1.13) (NADPH-GOGAT)
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01-DEC-1992 (Rel.
01-OCT-1994 (Rel.
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-I. SUBUNIT: ONE OF SEVEN SECRETORY PROTEINS (SECA-F & SECY)

COMPRISE THE PROKARYOTIC PROTEIN TRANSLOCATION APPARATUS.

-I. SUBCELLULAR LOCATION: Integral membrane protein.

-I. SIMILARITY: BELONGS TO THE SECY/SEC61-ALPHA FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biochemistry 31:3038-3043 (1992)
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Bacteria; Firmicutes; Bacillales; Geobacillus
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PS00756; SECY_2; PARTIAL.
transport; Translocation; Transmembrane.
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H., Barzu O., Gilles A.M.;
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2; Mismatches
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MEDLINE=93662802; PubMed=1435726;

MEDLINE=93662802; PubMed=1435726;

Tschauder S., Driessen A.J.M., Freudl R.;

Toloning and molecular characterization of the secy genes from "Cloning and molecular characterization of the secy genes from Bacillus licheniformis and Staphylococcus carnosus: comparative analysis of fine members of the Secy family.";

Mol. Gen. Genet. 235:147-152(1992)
                                                                                                                                                                                                                                                                                                                                                     TIGREAM6; TIGR00967; 3a0501
PROSITE; PS00755; SECY_1; 1
PROSITE; PS00756; SECY_2; 1
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COMPRISE THE PROKARYOTIC PROTEIN TRANSLOCATION APPARATUS
SUBCELLULAR LOCATION: Integral membrane protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FUNCTION: INVOLVED IN PROTEIN EXPORT. 1
TO ALLOW THE TRANSLOCATION OF PROTEINS
BY FORMING PART OF A CHANNEL.
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cranslocase secY subunit.
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EMBL; M31102; AAB5918.1; -.
EMBL; D00619; BAA00495.1; ALT_INIT.
EMBL; L47971; AAB06819.1; -.
EMBL; Z99104; CAB11912.1; -.
EMBL; Z99104; CAB11912.1; -.
PIR; S08629; BWBSSY.
PIR; S12683; S12683.
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Suh J.W., Boylan S.A., Oh S.H., Price C.W.;
"Genetic and transcriptional organization of the Bacillus subtilis
                       PRINTS; PR00303; SECYTRNLCASE.
TIGKERMS; TIGK00967; 3a05018007;
PROSITE; PS00755; SECY 1.1
PROSITE; PS00756; SECY_2; 1.
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Gene 169:17-23(1996)
-!- FUNCTION: INVOLV
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16-OCT-2001
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PRINTS; PR00303; SECYTF
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J. Biochem. 107:603-607(1990)
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Bacteria; Firmicutes;
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Protein transport; Translocation; Transmembrane; Complete proteome.
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Yoshikawa H., Doi R.H.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FUNCTION: INVOLVED IN PROTEIN EXPORT. INTERACTS WITH SECF TO ALLOW THE TRANSLOCATION OF PROTEINS ACROSS THE PLASMA BY FORMING PART OF A CHANNEL.
SUBJUNIT: ONE OF SEVEN SECRETORY PROTEINS (SECA-F & SECY) COMPRISE THE PROKARYOTIC PROTEIN RENASLOCATION APPARATUS. SUBJULLAR LOCATION: INTEGERAL Membrane protein. SUBJULLAR LOCATION: INTEGERAL Membrane protein.
                                                                                                                                                                                                                                                                                                                                                                                                             requires a license agreement (See an email to license@isb-sib.ch).
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Boylan S.A., Thomas S.M.,
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(Rel. 15, Last sequence)
(Rel. 40, Last annotation
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YP46_YEAST
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C. STRAIN-SCREEN, AB972;

X. MEDLINE-97313271; PubMed-9169875;

X. Bussey H., Storms R.K., Ahmed A., Albermann K., Allen E., Ansorge W.,

A. Araujo R., Aparicio A., Barrell B.G., Badcock K., Benes V.,

A. Chang E., Charcher C.M., Coster F., Davis K., Davis R.W.,

A. Chung E., Churcher C.M., Coster F., Davis K., Davis R.W.,

A. Dietrich F.S., Delius H., DiPaolo T., Dubois E., Duesterhoeft A.,

A. Dincan M., Floeth M., Fortin N., Friesen J.D., Fritz C., Goffeau A.,

A. Duncan M., Floeth M., Fortin N., Friesen J.D., Fritz C., Goffeau A.,

A. Hanil J., Hebling U., Heumann K., Hilbert H., Hillier L.,

A. Hunicke-Smith S., Hyman R., Johnston M., Kalman S., Kleine K.,

A. Hunicke-Smith S., Hyman R., Johnston M., Lin D., Louis E.J.,

A. Hanil J., Hebling U., Heumann K., Hilbert H., Lin A., Lin D., Louis E.J.,

A. Hanicke-Smith S., Hyman R., Johnston M., Lin D., Louis E.J.,

A. Hanicke-Smith S., Hyman R., Johnston M., Lin D., Louis E.J.,

A. Hanicke-Smith S., Hyman R., Johnston M., Lin D., Louis E.J.,

A. Hanicke-Smith S., Hyman R., Johnston M., Lin D., Louis E.J.,

A. Hanil J., Hebling U., Heumann K., Johnston M., Lin D., Louis E.J.,

A. Hanil J., Hebling U., Heumann R., Johnston M., Lin D., Louis E.J.,

A. Hanil J., Hebling U., Heumann R., Johnston M., Lin D., Louis E.J.,

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A. Hanil J., Hebling U., Heumann R., Johnston M., Lin D., Louis E.J.,

A. Hanil J., Hebling U., Heumann R., Johnston M., Lin D., Louis E.J.,

A. Hanil J., Hebling U., Heumann R., Johnston M., Scharfe M.,

A. Hanil J., Hebling U., Helmann R., Johnston M., Heller R., He
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16-OCT-2001 (Rel. 40, Last sequence update)
15-UIN-2002 (Rel. 41, Last annotation update)
Hypothetical protein YPL146C.
YPL146C OR LPI2C OR P2610
                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                      identifies a small nuclear RNA, a new new putative regulators."; Yeast 12:1483-1492(1996).
                                                                                                                                                                                                                                                                                                                                                                                    Purnelle B., Coster F., Goffeau A.;
"The sequence of 55 kb on the left arm of yeast chromosome XVI
                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=S288c / AB972;
MEDLINE=97103777; PubMed=8948103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Saccharomyces cerevisiae (Baker's yeast).
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16-0CT-2001
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EMBL; X96770; CAA55547.1; -.
EMBL; Z3502; CAA97850.1; -.
SGD; S0006067; YPLL46C.
HYPOCTHETICAL PROTECTION
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SEQUENCE
455 AA; 52555 MW;
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P15121; Q9BS21;
01-APR-1990 (Rel. 1
01-AUG-1992 (Rel. 2
15-JUN-2002 (Rel. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JUN-2002 (Rel. 23, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Aldose reductase (EC 1.1.1.21) (AR) (Aldehyde
AKRIBI OR ALDRI.
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Carey F., Markham A.F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; !
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and fc entities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
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-i- SIMILARITY; BELONGS TO THE GLTSCR2 FAMILY.
                          MEDLINE=90253609; PubMed=2111143; Grundmann U., Bohn H., Obermeier R.,
                                                                                                                                                                                                     "Cloning and sequence o
                                                                                                                                                                                                                                                                                                                                                          "Nucleotide sequence of cDNA for human aldose Nucleic Acids Res. 17:8368-8368(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=90045960; PubMed=2510130;
Graham A., Hedge P.J., Powell S.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bohren K.M., Bullock B., Wermuth B., Gabbay K.H.; "The aldo-keto reductase superfamily. cDNAs and deduced sequences of human aldehyde and aldose reductases."; J. Biol. Chem. 264:9547-9551(1989).
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MEDLINE=89255461; PubMed=2498333;
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                                                                                                       SEQUENCE FROM N.A.
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n U., Bohn H., Obermeier R., Amann E.;
and prokaryotic expression of a biologically active human
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"An unlikely sugar substrate site in the 1.65 A structure of
human aldose reductase holoenzyme implicated in diabetic
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"Structure of the human aldose reductase
J. Biol. Chem. 266:6872-6877(1991).
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Wilson D.K., Tarle I., Petrash J.M., Quiocho
"Refined 1.8-A structure of human aldose redu
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Jaquinod M., Potier N., Klarskov K.,
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J. Biol. Chem. 264:2912-29
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                                                   X-RAY CRYSTALLOGRAPHY (1.8 ANGSTROMS).
MEDLINE=98070237; PubMed=9405046;
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                                                                                                                                                                                           Biol. Chem.
                                                                                             inhibitor zopolrestat.";
Natl. Acad. Sci. U.S.A. 9
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V., Biellmann
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modified and this statement
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CATALYTIC ACTIVITY: Alditol + NAD(P)(+) = aldose + NAD(P)H.

STBUNIT: MONOMER.
                                                                                                                                                                                                                                                            DISEASE: IN DIABETES AND GALACTOSEMIA, INCREASED AR ACTIVITY LEADS TO HIGH LEVELS OF SORBITOL AND GALACTITIC, RESPECTIVELY, IN THE CELLS OF MANY TISSUES. ACCUMULATION OF SUGAR ALCOHOLS HAS BEEN SHOWN TO CAUSE OSMOTIC CATARACTS IN THE LENS. AR IS ALSO THOUGHT TO PLAY A KEY ROLE IN DIABETIC COMPLICATIONS OF THREE OTHER TARGET TISSUES, NAMELY, NERVE, KILDNEY AND RETURA.

SIMILARITY: BELONGS TO THE ALDO/KETO REDUCTASE FAMILY.
                                                                                              s SWISS-PROT entry is copyright. It is produced through a cween the Swiss Institute of Bioinformatics and the EMBL European Bioinformatics Institute. There are no restrict European Bioinformatics Institute. There are no restrict by non-profit institutions as long as its content is
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EMBL; X15414; CAA33460.1; -.
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EMBL; M34720; AAA35560.1; -.
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EMBL; M59863; AAA51712.1; -.
EMBL; M59863; AAA61712.1; -.
EMBL; BC000260; AAH00260.1; -.
EMBL; BC000380; AAH00260.1; -.
EMBL; BC000380; AAH00387.1; -.
EMBL; BC010391; AAH10391.1; -. MIM; 103880; -...
InterPro; IPR001395; Aldo/ket_red; I.
Pfam; pP00248; aldo_ket_red; I.
PRINTS; pR00069; ALDKETEDTASE.
ProDom; pD000288; Aldo/ket_red; I.
PROSITE; pS00062; ALDOKETO_REDUCTASE 2; 1.
PROSITE; PS00063; ALDOKETO_REDUCTASE 3; 1.
PROSITE; PS000798; ALDOKETO_REDUCTASE 1; 1.
PROSITE; PS00798; ALDOKETO_REDUCTASE 1; 1. Aarhus/Ghent-2DPAGE; A36436; A36510; A39763; IADS; 3; IABN; 3; IABN; 3; 2ACQ; 3; 2ACR; 3; 2ACS; 3; 1AZ2; 18-MAR-98. HGNC:381; AKR1B1 0; A36510.
33; A39763.
31-QXT-93.
31-JXN-94.
31-JUL-94.
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31-JUL-94. 20-JUL-95. 18-MAR-98. B33851. A31432. A36436. 18 48 48 141 141 13 13 13 1202; IEF HYDROGEN-BOND DONOR (BY SIMILARITY). L -> I (IN REF. 3). W -> R (IN REF. 5; AAH05387). IAE -> EAA (IN REF. 9). NADP ACETYLATION (POTENTIAL).

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16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
Muscarinic acetylcholine receptor M3 (Mm3 m
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                                                                                                                                                                                                                                                                                                                                                                  Andre C., Dos Santos G., Koulakoff A.;

"Cultured neurons from mouse brain reproduce the muscarinic receptor profile of their tissue of origin.";

Eur. J. Neurosci. 6:1691-1701(1994).

-i- FUNCTION: THE MUSCARINIC ACETYLCHOLINE RECEPTOR MEDIATES VARIOUS CELLULAR RESPONSES, INCLUDING INHIBITION OF ADENYLATE CYCLASE,

BREAKDOWN OF PHOSPHOINOSITIES & MODULATION OF POTASSIUM CHANNELS

THROUGH THE ACTION OF G PROTEINS. PRIMARY TRANSDUCING EFFECT IS PI
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MGD; MGI:88398; Chrm3.
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ACM3_N
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Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases.
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Mammalia; Eutheria; Rodentia;
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SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
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No.
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RESULT 14
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                                                                                            Bonner T.I., Young A.C., Brann M.R., "Cloning and expression of the human acetylcholine receptor genes."; Neuron 1:403-410(1988).
                           screening
                             MEDLINE=88077068; PubMed=3120722;
Braun T., Schofield P.R., Shivers B.D., Pritchett D.B.
"A novel subtype of muscarinic receptor identified by
screening.";
                                                                                                                                                                                 Bonner T.I., Buckley N.J., Young A.C., Brauldentification of a family of muscarinic
                                                                                                                                                                                                  SEQUENCE FROM N.A.
MEDLINE=87263421; PubMed=3037705;
                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus
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DOMAIN 1
TRANSMEM 67
SEQUENCE FROM N.A
                                                                 TISSUE=Brain;
                                                                          SEQUENCE FROM N.A.
                                                                                                                                   MEDLINE=90166521; PubMed=3272174;
                                                                                                                                            REVISION TO 184.
                                                                                                                                                                Science 237:527-532(1987)
                                                                                                                                                                                                                               NCBI_TaxID=10116;
                                                                                                                                                                                                                                                              Rattus norvegicus (Rat).
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PS50262; G_PROTEIN_RECEP_F1_2; 1.
ptic membrane; Ionic channel; Glycoprotein; Transmembrane;
pticn; Multigene family; G-protein coupled receptor.
ylation; Multigene family; G-protein coupled receptor.
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EXTRACELLULAR (POTENTIAL).
                  Biophys.
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nic acetylcholine
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PIR; ....
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Pfam; PF00001; 7tm 1; 1.

Pfam; PF00001; 7tm 1; 1.

PRINTS; PR00237; GFORRHODOPSN.

PRINTS; PS00237; G PROTEIN RECEP F1 2; 1.

PROSITE; PS00237; G PROTEIN RECEP F1 
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EMBL; M16408; AAA40662.1; ALT_SEQ.

EMBL; M18089; AAA40659.1; -.

EMBL; M62826; AAA41553.1; -.

EMBL; AB017656; BAA36839.1; -.

EMBL; AB017656; BAA36839.1; -.

PIR; B29514; B29514.

PIR; A29476; A29476.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      muscarinic receptor point mutants.";
J. Biol. Chem. 267:19313-19319 (1992).
J. Biol. Chem. 267:19313-19319 (1992).
I- FUNCTION: THE MUSCARINIC ACETYLCHOLINE RECEPTOR MEDIATES VARIOUS
-I- FUNCTION: THE MUSCARINIC ACETYLCHOLINE RECEPTOR MEDIATES VARIOUS
-I- FUNCTION: THE MUSCARINIC ACETYLCHOLINE INCLUDING INHIBITION OF ADENYLATE CYCLASE,
BREAKDOWN OF PHOSPHOINOSITIDES & MODULATION OF POTASSIUM CHANNELS
THROUGH THE ACTION OF G PROTEINS. PRIMARY TRANSDUCING EFFECT IS PI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE-92037535; PubMed=1657592;
Wess J., Gdula D., Brann M.R.;
"Sitce-directed mutagenesis of the m3 muscarinic receptor:
identification of a series of threonine and tyrosine resi
involved in agonist but not antagonist binding.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Furuta M., Ohya S., It "Molecular cloning of iris.";
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Furuta M., Ohya S., Imaizumi Y., Wat
"Molecular cloning of m3 muscarinic
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SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS
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01-NOV-1995 (Rel. 3
01-NOV-1995 (Rel. 3
15-JUL 1998 (Rel. 3
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Sui A.-L., Chou W.-Y., Kao L.-S.;
Submitted (XXX-1994) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: THE MUSCARINIC ACETYLCHOLINE RECEPTOR MEDIATES VARIOUS
-ICELLULAR RESPONSES, INCLUDING INHEBITION OF ADENYLATE CYCLASE,
BREAKDOWN OF PHOSPHOINOSITIDES & MODULATION OF POTASSIUM CHANNELS
THROUGH THE ACTION OF G PROTEINS. PRIMARY TRANSDUCING EFFECT IS PI
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EMBL; L27103; AAA30653.1;
InterPro; IPR000276; GPCR_Rhodpsn.
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PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
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acetylcholine receptor M3.
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MEDLINE=92299690; PubMed=1376732;

Hardwick C., Hoare K., Owens R., Hohn H.P., Hook M., Moore D.,

Cripps V., Austen L., Nance D.M., Turley E.A.;

"Molecular cloning of a novel hyaluronan receptor that mediates tumor
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Hardwick C., Hoare K., Owens R., Hohn H.P., Hook M., Moore D.,
Cripps V., Austen L., Nance D.M., Turley E.A.;
"Correction. Molecular cloning of a novel hyaluronan receptor that
mediates tumor cell motility.";
J. Cell Biol. 118:753-753(1992).
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J. Cell Biol. 11
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   423
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KLRSQLVKRK 432
                                               KLKSOLVKRK 10
                                                                                                                                         Similarity 9; Conserv
                                                                                                                                                                                                                                                                               476 AA;
                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                               55486 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chordata; Craniata; Vertebrata; Euteleostomi; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                          93.5%;
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Last annotation update)
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                                                                                                                                         Score 43; DB
Pred. No. 2.4;
1; Mismatches
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                                                                                                                                                                                                                                                                                   62DCA82045EE0CB2 CRC64;
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                                                                                                                                                                                                          DB 11; Length 476;
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Best Local Similarity
Matches 8; Conserv
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01-NOV-1999
01-JUN-2001
                                                                                                                                                                                                                                                                  Receptor.
SEQUENCE
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SEQUENCE
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01-MAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
5imilar to hyaluronan mediated motility receptor (RIMBL musculus (Mouse).
                                                                                                                                                                                                                                                                                                  TISSUE-PHEOCHROMOCYTOMA;
Lynn B.D., Li X., Cattini P.A., Nagy J.I.;
"Sequence, protein expression and erk associat
RHAMM in PC12 cells.";
Submitted (JAN-2001) to the EMBL/GenBank/DDBJ
EMBL; AF133037; AAD24473.1; -.
EMBL; AF336825; AAKZ1904.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN-SPRAGUE DAWLEY;
Li X., Lynn B., Nagy J.
"RHAMM cDNA from rat bu
Submitted (MAR-1999) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Strausberg R.;
Submitted (JAN-2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hyaluronan
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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99 (TrEMBLrel. 12, 1
01 (TrEMBLrel. 17, 1
n receptor RHAMM.
                                                                                                                                                                                                                                                                  713 AA;
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                                                                                                                                                                                                                                                                82395 MW;
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Last sequence
Last anno
                                                                                                                                                                          Score 39; DB
Pred. No. 21;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL/GenBank/DDBJ databases.
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Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Rat
                                                                                                                                                                                                                                                                  2FE310D5759C6CB3 CRC64;
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Q942A1;
01-DEC-2001
01-DEC-2001
01-MAR-2002
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O9FRZ1;
01-MAR-2001 (TrEMBLrel 16,
01-MAR-2001 (TrEMBLrel 16,
01-MAR-2002 (TrEMBLrel 20,
                                                                                                                                   STRAIN-GOLDEN CROSS BANTAM T51;
Sakakibara H., Sugiyama T.;
"Cloning and characterization of maize response regulators.";
Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AB042261; BAB20579.1; -.
InterPro, IPR001789; Response reg.
Pfam; PF00072; response_reg; 1.
                                                                                                 ProDom; PD000039; Res
SMART; SM00448; REC;
Phosphorylation; Sen
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Spermatophyta; Magnoliophyta;
Panicoideae; Andropogoneae; Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (MAY-2001) to the EMBL/GenBank/DDBJ EMBL, AP003683; BAB64697.1; -.
INTERPO, IPR001789; Response reg.
Pfam; PF00072; response reg; T.
ProDom; PD000039; Response reg; 1.
Phosphorylation; Sensory transduction.
Phosphorylation; Sensory transduction.
SEQUENCE 232 AA; 24863 MW; 5454CA63CF2EF95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sasaki T., Matsumoto T., Yamamoto K.;
"Oryza sativa nipponbare(GA3) genomic
clone:P0431G6.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Viridiplantae; Streptophyta; Er
Spermatophyta; Magnoliophyta; Liliopsida;
Ehrhartoideae; Oryzeae; Oryza.
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                                                                                         SEQUENCE
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                     KLKSQLVKRK 10
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KLKSHLLKRK 157
                                          8; Conserv
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1 (TrEMBLrel. 19,
2 (TrEMBLrel. 20,
                                                                                       235 AA;
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                                           Conservative
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                                                                                                    Sensory transduction.
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                                                     82.6%;
80.0%;
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yta; Liliopsida;
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                                          Pred. No. 12;
L; Mismatches
                                                                 Score 38;
                                                                                         A53C205400141DCE CRC64;
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a; Poales; Poaceae; PACC cl
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RESULT 7

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STRAIN=ILI403;

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MEDLINE=21235186; PubMed=11337471;

Malper S., Jaillon O., Ma Weissenbach J., Ehrlich S.D., Sorokin A.;

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EMBL; AB006263; AAK04355.1; -.
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SEQUENCE FROM N.A.
STRAIN-CV. GOLDEN CROSS BANTAM T51;
Sakakibara H., Sugiyama T.;
Sakakibara and characterization of maize response regulators.";
"Cloning and characterization of maize response regulators.";
Submitted (Apr-2000) to the EMBL/GenBank/DDBJ databases.
Submitted (Apr-2001);
EMBL; AB042267; BAB20580.1;
EMBL; AB042267; BAB20580.1;
InterPro; IPR001789; Response reg.
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Pfam; PF00072; response reg; I.
ProDom; PD000039; Response reg; 1.
SMART; SM00448; REC; 1.
Phosphorylation; Sensory transduction.
SEQUENCE 236 AA; 25454 MW; 5P8CEC4
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Submitted (CCT-2001) to the

EMBL; U97194; AAL02460.1; -.

Hypotherical protein.

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TT.T., Kemp K., Scheet P.;

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"Direct Submission.";
Submitted (SEP-2001) to the EMBL; U97194; AAB52452.1; -
Hypothetical protein.
SEQUENCE 336 AA; 39659 M
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01-JUN-2002 (
01-JUN-2002 (
Receptor for
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NON_TER
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SEQUENCE
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  Schoenfelder M., Einspanier R.; "Exgression of HAS-system during oocyte maturation Submitted (MAR-2002) to the EMBL/GenBank/DDBJ datab EMBL; AJ439694; CAD29126.1; -.
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2002 (TrEMBLrel. 21, Last annotation update)
101-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Hyaluronic acid-mediated motility receptor (Fragment)
Ovis aries (Sheep).
                                                                                                                                                                  Eukaryota; Metazoa; Ch
Mammalia; Eutheria; Ce
Bovidae; Bovinae; Bos.
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NCBI_TaxID=9940,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Receptor for hyaluronic
                                                                                     rissue=ovary;
                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                             NCBI_TaxID=9913;
                                                                                                                                                                                                                                Bos taurus
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Cetartiodactyla; Ruminantia;
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Matches 7
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Q9Z607;
Q9Z607;
01-MAY-1999
01-MAY-1999
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Q9S689;
01-MAY-2000
01-MAY-2000
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SEQUENCE
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SEQUENCE FROM N.A.
STRAIN-MBA TYPE 8;
MEDLINE=99045873; PubMed=9828433;
MEDLINE=99045873; PubMed=9828433;
SEQUENCE FROM N.A.
STRAIN=MBA TYPE X;
MEDLINE=98411422; PubMed=9738062;
                                                                             Bacteria; Firmicu
Mycoplasmataceae;
NCBI_TaxID=2130;
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STRAIN=MBA TYPE 8;
MEDLINE=98411422;
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Multiple banded antigen
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Clin. Microbiol. 36
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L; AF108876; AAD19955.1;
_TER 68 68
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ment.";
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Ureaplasma urealyticum (Ureaplasma urealyticum biotype 2).
Bacteria; Firmicutes; Bacillus/Clostridium group; Mollicutes;
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Bacteria; Firmicutes; Bacillus/Clostridium group; Mollicutes;
Mycoplasmataceae; Ureaplasma.
                                   (TrEMBLrel. 10, Created)
(TrEMBLrel. 10, Last sequence update)
(TrEMBLrel. 16, Last annotation update)
nded antigen (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nested PCR, and random nd typing of Ureaplasma
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Pred. No. 15;
0; Mismatches
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Pred. No.
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                                                  Q9ZB37 PRELITATION...,
Q9ZB37 PRELITATION...,
Q9ZB37;
Q1-MAY-1999 (TrEMBLrel. 10, Last sequence update)
Q1-MAY-1999 (TrEMBLrel. 16, Last annotation update)
Q1-MAR-2001 (TrEMBLrel. 16, Last annotation update)
Multiple banded antigen (Fragment).
Ureaplasma urealyticum (Ureaplasma urealyticum biotype 2).
Bacteria, Firmicutes, Bacillus/Clostridium group; Mollicutes;
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J. Clin. Microbia
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Q9ZB38;
Q1-MAY-1999 (TrEMBLrel. 10, Created)
O1-MAY-1999 (TrEMBLrel. 10, Last sequence update)
O1-MAR-2001 (TrEMBLrel. 16, Last annotation update)
O1-MAR-2001 (TrEMBLrel. 16, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zheng X., Teng L.-J., Watson H.L., Glass J.I., Cassell G.H "Sequence analysis of the major servour specific antigen (I genes from different servours of Ureaplasma urealyticum."; Submitted (MAR-1996) to the EMBL/GenBank/DDBJ databases.
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STRAIN-MBA TYPE X;
MEDLINE-99045873; PubMed-9828433;
Knox C.L., Giffard P., Timms P.;
  Zheng X., Teng L.-J., "Sequence analysis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Multiple banded antigen (Fragment)
Ureaplasma urealyticum (Ureaplasma urealyticum biotype 2)
Bacteria; Firmicutes; Bacillus/Clostridium group; Mollicutes;
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fragment.";
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ER 186 186
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AF108877; AAD19956.1; -.
ER 68 68
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Watson H.L., Glass J.I., Cassell G. the major serovar specific antigen
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Best Local Similarity
Matches 8; Conserv
                                                                                               genes from different serovars of Ureaplasma urealyticum."; Submitted (MAR-1996) to the EMBL/GenBank/DDBJ databases. EMBL, U50461; AAD00077.1; -.
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KLSSQLVKSK
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Search completed: February 8, 2003, 10:42:44 Job time: 23.6667 secs

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2: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:*
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AAW56543	AAW17884	AAW56544	AAW17885	ABG28286	ABB68353	ABG00402	ABG12556	ABG19127	ABG19121	ABB71860	ABG19123	ABG19124	ABG00387	ABG12563	ABG00388	ABG19125	ABG19122	AAB96656	AAE24061	ABB95331	AAM01226	AAU69871	ABB60907	ABB83356	AAW69357	ABB53563	AAB48053	AAU11442	ABG60850	AAR46550	AAU11438	AAW39168	AAU11437	ABG60843
Toxin TcaB, encode	Photorhabdus lumin	Toxin TcaBi, encod	Photorhabdus lumin	Novel human diagno	Drosophila melanog	human		human	Ħ	Drosophila melanog	Novel human diagno	Novel human diagno	Novel human diagno	Novel human diagno	Novel human diagno	human		Putative P. abyssi	prostat		Human prostate-spe	Human prostate cDN	Drosophila melanog	Murine TSAP6 bindi	Rat lens aldose re	Lactococcus lactis	Signal transductio	Hyaluronic acid-bi	Cellular response	Hyaluronan recepto	Rat hyaluronic aci	Rat RHAMM binding	Mouse hyaluronic a	Mouse receptor for

ALIGNMENTS

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13-AUG-2002
                                                                      ABG60827;
                                                                                                      ABG60827 standard; Peptide; 10
Hyalauronan (HA) binding peptide #10
                                   (first entry)
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Tissue disorder; response-to-injury process; cell proliferating; hyaluronic acid; HA; receptor for hyalauronan-mediated motility; RHAMM; inflammatory neurological disorder; Parkinson's disease; Alzheimer's disease; arthritis; multiple sclerosis; gastritis; nephritis; inflammatory dermatosis; pooriasis; inflammatory bowel disease; stenosis; restenosis; cancer; kidney fibrosis; inflammatory lung disease; emphysema; asthma; cystic fibrosis; obesity; obesity related disease; nupus; cardiovascular disease; athroseclerosis; wound; scar; diabetes; tissue transplantation; stroke; inflammatory response; fibrotic response; medical implant; Acquired immunodeficiency syndrome; AIDS; hepatitis; myocardial fibrosis; hepatic fibrosis; chronic cystitis; acute mastitis; 05-OCT-2000; 2000WO-IB01534. 05-OCT-2000; 2000WO-IB01534. 11-APR-2002 WO200228415-A1. Synthetic. septic shock; thyroiditis; retinopathy.

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RESULT 2
AAA46551
ID AAA4
XX AAA4
AC AAA4
AC AAA4
AC AAA4
XX DT 05-A
XX Hyal
DE Hyal
KW Heal
KW heal
KW birt
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention describes a method of treating a tissue disorder associated CC with response-to-injury process or proliferating cells in a patient, CC comprising administering a polypeptide (I) which binds hyaluronic acid (HA), an antibody which binds one of domains D1-D5 of Receptor for CC hyalauronan-mediated motility (RHAMM), a polypeptide fragment encoding CC any of D1-D5 of RHAMM, or a vector which expresses antisense RHAMM, CC antibodies or a polypeptide fragment. The method is useful for treating a CC disease, Alzheimer's disease, arthritis including rheumatoid arthritis, constantially selected in the method is useful for treating a costeoarthritis, multiple sclerosis, inflammatory dermatosis (psoriasis). CC inflammatory bowel disease, stenosis or restenosis, cancer, kidney CC fibrosis, inflammatory lung disease (e.g. emphysema, asthma, cystic CC fibrosis, obesity or obesity related diseases, lupus, cardiovascular CC disease (e.g. atherosolerosis), and wound especially surgical excision CC adhesions, to prevent scar and also for treating or preventing diabetes CC mellitus. The method is also useful for treating tissue transplantation (e.g. skin grafts), stroke, inflammatory responses or fibrotic response associated with medical implants such as hIDS, myocardial and hepatic CC fibrosis, chronic cystitis, acute mastitis, saguritis, nephritis, compressents a hyalauronan (RA) binding peptide used in the method of treating a tissue disorder described in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
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                                                                                                                                                                        healing; diagnosis; treatment; cell locomotion; tumour invasion; birth defects; inflammatory disorder; Alzheimer's disease; dementia; Parkinson's diseases; Hutington's disease; AIDS; diabetes; auto; immune diseases; corneal dysplasia; hypertrophy; surgery; burns; strokes; multiple sclerosis; depression; schizophrenia; CNJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating tissue disorder associated with response-to-injury process or proliferating cells in mammals, e.g. fibrosis, inflammation, by administering a compound that alters activity of transition molecules within a cell
                                      13-APR-1993;
                                                                                                                                                        contraception;
                                                                                                                                                                                                                                                                          Hyaluronan binding protein; HA; RHAMM; mediated motility; wound;
                                                                                                                                                                                                                                                                                                                 Hyaluronan receptor binding
                                                                                                                                                                                                                                                                                                                                                         05-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                AAR46551;
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                                      93WO-CA00158
                                                                                                                                                          vitro
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Pred. No. 0.026;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RHAMM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    wound healing, in response to growth factors and in chemotaxis by white blood cells. HA may be used for diagnosis and treatment of diseases involving cell locomotion, e.g. tumour invasion, birth defects, acute and chronic inflammatory disorders, Alzheimer; s and other forms of dementia, AIDS, diabetes, autoimmune diseases, corne dysplasias and hypertrophies, burns, surgical incisions and adhesic strokes, multiple sclerosis, depression/schizophrenia related to neuronal growth and pain states involving nerve sprouting; also in
     Isolated human receptor to develop products for
                                                                                   WPI; 1997-512715/47.
                                                                                                                                 Entwistle J,
                                                                                                                                                                                          (MANI-)
                                                                                                                                                                                                                                                                                                                                                                            16-OCT-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                            WO9738098-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequence is that of a binding motif fragment of the h receptor (HARC). HARC is down regulated in normal cells expressed in situations where cell motility is desired, e
                                                                                                                                                                                                                                                                     10-APR-1996;
                                                                                                                                                                                                                                                                                                                      10-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hyaluronan receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse RHAMM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW39164 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and spinal cord regeneration,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 18; Page 57; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA encoding hyaluronan receptor - used to produce proteins and antibodies for alteration of cell locomotion
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammatory disorder; autoimmune disease; diagnosis; prognosis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      onan receptor; receptor for hyaluronic acid mediated motility;
glycosaminoglycan; binding domain; mouse; oncogene; treatment
factor; cell locomotion disorder; dementia; detection;
                                                                                                                                                                                       MANITOBA CANCER UNIV MANITOBA.
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                                                                                                                                 Turley
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                                                                                                                                                                                                                                                                     96GB-0007441
                                                                                                                                                                                                                                                                                                                         97WO-CA00240
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for hyaluronic acid mediated motility - treating e.g. tumours, inflammatory
                                                                                                                                                                                                                  TREATMENT & RES FOUND
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Pred. No. 0.097;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               consensus motif
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RHAMM; inflammatory neurological disorder; Parkinson's disease; Alzheimer's disease; arthritis; multiple sclerosis; gastritis; nephritis; inflammatory bowel disease; inflammatory dermatosis; psoriasis; inflammatory bowel disease; stenosis; restenosis; cancer; kidney fibrosis; inflammatory lung disease; emphysema; asthma; cystic fibrosis; obesity; obesity related disease; lupus; cardiovascular disease; atherosclerosis; wound; scar; diabetes; tissue transplantation; stroke; inflammatory response; fibrotic response; medical implant; Acquired immunodeficiency syndrome; ALDS; hepatitis; myocardial fibrosis; hepatic fibrosis; chronic cystitis; acute mastitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                be used in e.g. CNS and spinal cord regeneration, contraception and in vitro fertilisation and embryo development. The products can also be
                                                WPI; 2002-435298/46
                                                                                                                                                                                                                                                                                                                                                                                      05-OCT-2000; 2000WO-IB01534.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tissue disorder,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cellular response to injury associated peptide #3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABG60822;
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                                                                                                                                 Turley EA,
                                                                                                                                                                                                                                                                                                 05-OCT-2000; 2000WO-IB01534.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hyaluronic acid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-AUG-2002
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nes 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 KLRSQLVKRK 10
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                                                                                                                                                                                                                 TRANSITION THERAPEUTICS & DIAGNOSTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         dementia, AIDS, diabetes and auto-immune diseases
                                                                                                                            Cruz TF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               thyroiditis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        response-to-injury process; cell proliferating; HA; receptor for hyalauronan-mediated motility; ory neurological disorder; Parkinson's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnosis and prognosis.
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Pred. No. 0.097;
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DB 23;

Length 11;

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fibrosis), obesity or obesity related diseases, lupus, cardiovascular disease (e.g. atherosclerosis), and wound especially surgical excision adhesions, to prevent scar and also for treating or preventing diabetes mellitus. The method is also useful for treating tissue transplantation (e.g. skin grafts), stroke, inflammatory responses or fibrotic response associated with medical implants such as hip implants, vascular wraps an catheters), inflammatory diseases such as AIDS, myocardial and hepatic fibrosis, chronic cystitis, acute mastitis, gastritis, nephritis, hepatitis, septic shock, thyroiditis, and retinopathy. This sequence represents a peptide associated with the method of treating tissue disorders described in the invention.
                                                                                                                                                                                                                                                                                                                                                                                        (HA), an antibody which binds one of domains D1-D5 of Receptor for hyalauronan-mediated motility (RHAMM), a polypeptide fragment encoding any of D1-D5 of RHAMM, or a vector which expresses antisense RHAMM, antibodies or a polypeptide fragment. The method is useful for treating a patient with an inflammatory neurological disorder such as Parkinson's disease, Alzheimer's disease, arthritis including rheumatoid arthritis, osteoarthritis multiple sclerosis, inflammatory bowel disease, stenosis or restenosis, cancer, kidney inflammatory bowel disease, stenosis or restenosis, cancer, kidney inflammatory lung disease (e.g. emphysema, asthma, cystic fibrosis, inflammatory lung disease (e.g. emphysema, asthma, cystic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Treating tissue disorder associated with response-to-injury process or proliferating cells in mammals, e.g. fibrosis, inflammation, by administering a compound that alters activity of transition molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                with response-to-injury process or proliferating cells in a patient, comprising administering a polypeptide (I) which binds hyaluronic acid (HA), an antibody which binds one of domains D1-D5 of Receptor for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 199;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR99677 standard; Protein; 32
                                                                                   10-JUL-1996
                                                                                                                                                        EP721012-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-0CT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        breast cancer; therapy; dominant suppressor mutant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Domain
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9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                             /label= HA_binding_domain_II
/note= "corresponds to amino
RHAMM"
                                                                                                                                                                                                                                                                                                                                                                                                                /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label= HA_binding_domain_1
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                                                                                                                                                                                                                                                                                                                                                                                         "corresponds to amino acids 532-542 of RHAMM"
                                                                                                                                                                                                                                                                                                                                                                                 RHAMM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 43; DB Pred. No. 0.11
1; Mismatches
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RESULT 6
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Best Local
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                                                                                                                                                        Tissue disorder; response-to-injury process; cell proliferating; hyaluronic acid; HA; receptor for hyalaurona-mediated mottlity; RHAMM; inflammatory neurological disorder; Parkinson's disease; Alzheimer's disease; arthritts; multiple sclerosis; gastritts; nephritis; inflammatory dermatosis; psoriasis; inflammatory bowel disease; stenosis; restenosis; cancer; kidney fibrosis; inflammatory lung disease; emphysema; asthma; cystic fibrosis; obesity; obesity related disease; lupus; cardiovascular disease; atherosclerosis; wound; scar; diabetes; tissue transplantation; stroke; inflammatory response; fibrotic response; medical implant, Acquired immunodeficiency syndrome; AIDS; hepatitis; myocardial fibrosis; hepatic fibrosis; chronic cystitis; acute mastitis; septic shock; thyroiditis; retinopathy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The hyaluronan (HA) binding domains (AAR99677) of receptor for hyaluronan caid-mediated motility protein, RHAMM 1 (AAR99673), were mutated (see also AAR99678) in order to determine the importance of RHAMM for maintenance of the transformed phenotype of H-ras-transformed fibrosarcomas. Loss of HA binding was confirmed for the mutated RHAMM fusion protein in a transblot assay. Reversion of H-ras-transformation with the mutated RHAMM was demonstrated. Dominant suppressor mutants of the RHAMM gene may be used in the treatment or prevention of disorders characterised by RHAMM gene
05-OCT-2000; 2000WO-IB01534
                              05-OCT-2000; 2000WO-IB01534.
                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                                                                                                                    Receptor for hyalauronan-mediated motility binding peptide #2.
                                                                                                                                                                                                                                                                                                                                                                                                         13-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABG60846 standard; Peptide; 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             overexpression, e.g. breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; Fig 18A; 117pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Receptor for hyaluronic acid-mediated motility protein, and DNA encoding it - useful to treat or prevent diseases associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; AAT34524.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Entwistle J, Turley EA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-OCT-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 KLKSQLVKRK 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          receptor, e.g. breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KLRSQLVKRK 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT34500;
AAT23423;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ₽
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  . 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 43;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ₿
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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                                                                                                                                                                                   CC with response to injury process or proliferating cells in a patient, CC comprising administering a polypeptide (I) which binds hyaluronic acid CC (HA), an antibody which binds one of domains D1-D5 of Receptor for CC hyalauronan-mediated motility (RHAMM), a polypeptide fragment encoding CC any of D1-D5 of RHAMM, or a vector which expresses antisense RHAMM, CC antibodies or a polypeptide fragment. The method is useful for treating a CC patient with an inflammatory neurological disorder such as Parkinson's CC disease, Alzheimer's disease, arthritis including rheumatoid arthritis, CC disease, bowel disease, stenosis, inflammatory dermatosis (psoriasis), inflammatory bowel disease, stenosis or restenosis, cancer, kidney CC fibrosis, inflammatory lung disease (e.g. emphysema, asthma, cystic CC fibrosis, bobesity or obseity related diseases, lupus, cardiovascular CC disease (e.g. atherosclerosis), and wound especially surgical excision CC adhesions, to prevent scar and also for treating or preventing diabetes CC mellitus. The method is also useful for treating tissue transplantation CC (e.g. skin grafts), stroke, inflammatory responses or fibrotic response CC asheters), inflammatory diseases such as hip implants, vascular wraps and CC catheters), inflammatory diseases such as hips, myocardial and hepatic CC fibrosis, chronic cystitis, acute mastitis, gastritis, nephritis, represents a receptor for hyalauronan-mediated motility binding peptide creating a trease disorder described in the method of treating a tissue disorder described in the
                                                                                         Matches
                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Treating tissue disorder associated with response-to-injury process or proliferating cells in mammals, e.g. fibrosis, inflammation, by administering a compound that alters activity of transition molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 11; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               within a cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (TRAN-) TRANSITION THERAPEUTICS & DIAGNOSTICS
    23
                                            1 KLKSQLVKRK 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention describes a method of treating a tissue disorder associated
KLRSQLVKRK 32
                                                                                                                                                                                                                                                  in the method of treating a tissue disorder described in the
                                                                                         9
                                                                                                                                                                                   32
                                                                                       Conservative
                                                                                                                                                                                   B
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                                                                                                              93.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                215pp; English
                                                                                         1; Mismatches
                                                                                                              Score 43; I
Pred. No. 0.
                                                                                                                                    BB
                                                                                                                                      23;
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                                                                                                                                 Length 32
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                                                                                         Gaps
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PF XXX
                                                                                                                                                                         Hyaluronan binding protein; HA; RHAMM; mediated motility; wound; healing; diagnosis; treatment; cell locomotion; tumour invasion; birth defects; inflammatory disorder; Alzheimer's disease; demen
13-APR-1993;
                                  28-OCT-1993
                                                                                                      strokes; multiple sclerosis; depression; contraception; in vitro fertilisation; en
                                                                                                                  birth defects; inflammatory disorder; Alzheimer's disease; dementia; Parkinson's diseases; Huntington's disease; AIDS; diabetes; auto; immune diseases; corneal dysplasia; hypertrophy; surgery; burns; strokes; multiple sclerosis; depression; schizophrenia; CNJ;
                                                                                                                                                                                                                                                                                     05-APR-1994 (first entry)
                                                                                                                                                                                                                                                                                                                                                       AAR43563 standard; Protein; 476 AA.
                                                                                                                                                                                                                                                  Hyaluronan
                                                                     WO9321312-A
                                                                                                                                                                                                                                               receptor
93WO-CA00158
                                                                                                        embryo development
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expressed in situations where cell motility is desired, e.g. in wound healing, in response to growth factors and in chemotaxis by white blood cells. HA may be used for diagnosis and treatment of diseases involving cell locomotion, e.g. tumour invasion, birth defects, acute and chronic inflammatory disorders, Alzheimer; s and other forms of dementia, AIDS, diabetes, autolimmune diseases, corneal dysplasias and hypertrophies, burns, surgical inclisions and adhesions, strokes, multiple sclerosis, depression/schizophrenia related to neuronal growth and pain states involving nerve sprouting; also in CNJ and spinal cord regeneration, contraception, in vitro fertilisation and embroo demonstrates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The sequence is that encoded by a cDNA clone encoding the hyaluronan receptor (HARC). The sequence was obtd. by screening a 3T3 library in lambda gt11 with antibodies to ARC. A clone of 1.9 kb was obtained and used to rescreen the library to obtain the full length, 2.9 kb clone. HA is down regulated in stationary normal cells and is only expressed in situations where cell motility is desired, e.g. in
                                                                                                                                                                                            BuM
                                                                                                                                                                                                                                    RHAMM 1; receptor for hyaluronic acid hyaluronan receptor; cell locomotion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA encoding hyaluronan receptor - used to produce antibodies for alteration of cell locomotion
                                                                                                                   Modified-site
                                                                                                                                                                                                                                                                                 Receptor for hyaluronic acid mediated motility RHAMM 1.
                                                                                                                                                                                                                                                                                                                                             AAR99673;
                                                                                                                                                                                                                                                                                                                                                                     AAR99673 standard; Protein; 606 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 7; Fig 23; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-APR-1992;
 Modified-site
                              Modified-site
                                                                                      Modified-site
                                                                                                                                                                                                                         breast cancer; therapy.
                                                                                                                                                                                                                                                                                                              10-OCT-1996
                                                          Modified-site
                                                                                                                                               Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                    423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MANI-) MANITOBA CANCER TREATMENT & RES FOUND. (UYMA-) UNIV MANITOBA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
nes 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 also AAR46548-51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1993-351722/44.
DB; AAQ51212.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KLKSQLVKRK 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  development
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     476 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92GB-0007949
                                                                                                                           Location/Qualifiers
91..93
/label= N-glycosylation_site
/label= N-glycosylation_site
342..344
                              321..323
                                                            /label= N-glycosylation_site
300..302
                                                                                      /label= N-glycosylation_site
279..281
                                         /label= N-glycosylation_site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93.5%;
                                                                                                                   .260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 43;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         red. No. 4.2;
Mismatches
                                                                                                                                                                                                                                  mediated motility;
cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14; Length 476;
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ARESULT 9
AAW39166
ID AAW3
XX AAW3
AC AAW3
XX AAW3
XX AAW3
XX AAW3
XX MOU!
XX MOU!
XX HyA.
XX HYA.
XX RHA!
XX GRA!
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                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                     Query Match
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N-PSDB; AAT34500;
N-PSDB; AAT23423;
N-PSDB; AAT34524.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB;
N-PSDB;
N-PSDB;
                                                                                                                                                                                                                                                                                                                                     RHAMM 1 (AAR99673), or Receptor for Hyaluronic Acid Mediated Motility, is a hyaluronan receptor protein which is involved in cell locomotion or motility and cell proliferation and transformation. Its amino acid sequence was deduced from a cDNA clone (AAT34409) obtd. from murine 3T3 cells and a genomic sequence (AAT34500) from a mouse fibroblast genomic library. 2 Alternative mRNAs for RHAMM 1 were found, RHAMM 1A (see also AAT34523) and RHAMM 1B (AAT34524), that had identical translated portions. Increased expression of RHAMM 1 protein can be used to suppress or control a tumour by modulating the interaction of cell-associated RHAMM with its ligand.
Hyaluronan receptor; receptor for hyaluronic acid mediated motility; RHAMM; glycosaminoglycan; binding domain; mouse; oncogene; treatment growth factor; cell locomotion disorder; dementia; detection;
                                                      Mouse RHAMM protein.
                                                                                 27-APR-1998
                                                                                                                                    AAW39166 standard; Protein; 630
                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; Page 40-42; 117pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the receptor, e.g. breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Receptor for hyaluronic acid-mediated motility protein, and DN encoding it - useful to treat or prevent diseases associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1996-310997/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Entwistle J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-OCT-1994;
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                                                                                                                                                                                                       553
                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                              1 KLKSOLVKRK 10
                                                                                                                                                                                                                                                         Similarity
9; Conserv
                                                                                                                                                                                                                                                                                                                 909
                                                                                                                                                                                                                                                           Conservative
                                                                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Turley
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373..375
/label= N-glycosylation_site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            553..562
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594..596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /label= N-glycosylation_site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /label= Hyaluronan_binding_domain-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /label= N-glycosylation_site
                                                                                                                                                                                                                                                                       93.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        _N-glycosylation_site
                                                                                                                                                                                                                                                          Score 43; DB
Pred. No. 5.3;
1; Mismatches
                                                                                                                                    B
                                                                                                                                                                                                                                                                        DB 17;
5.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FOUND
                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                     Length 606;
                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and DNA
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RESULT 10
AAR99675
ID AAR99
XX
AC AAR99
XX
AC AAR99
XX
AC AAR99
XX
AC AAR99
XX
XX
AC AAR99
XX
XX
XX
DE RHAMM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC This sequence represents the mouse hyaluronan receptor which is is also CC known as the receptor for hyaluronic acid mediated motility (RHAMM). CC Hyaluronan is a large glycosaminoglycan that is ubiquitous in the CC extracellular matrix and whose synthesis has been linked to cell CC migration, growth and transformation. It interacts with cell surfaces via specific protein receptors, e.g. RHAMM, that mediate many biological CC effects. The RHAMM/Hyaluronic acid interaction is involved in CC oncogene-and growth factor-mediated cell locomotion. The products can be CC used in the treatment of disorders involving cell locomotion, e.g. tumour CC invasion, birth defects, acute and chronic inflammatory disorders, CC Alzheimer's and other forms of dementia, including parkinson's and CC Huntington's diseases, AIDS, diabetes, autoimmune diseases, corneal CC dysplasia and hypertrophies, burns, surgical incisions and adhesions, CC strokes and multiple sclerosis. They can also be used in e.g. CNS and CC embryo development. The products can also be used in detection, diagnosis CC and prognosis.
                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
Region
                       RHAMM 1-2a isoform
                                                                   10-OCT-1996
                                                                                                               AAR99675
                                                                                                                                                       AAR99675 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 46; 66pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Isolated human receptor for hyaluronic acid mediated motility to develop products for treating e.g. tumours, inflammatory disorders, dementia, AIDS, diabetes and auto-immune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Entwistle J,
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2
8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-APR-1996;
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                                                                                                                                                                                                                                                                         577
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                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                   ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prognosis.
                                                                                                                                                                                                                                                                                                                KLKSQLVKRK 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1997-512715/47
                                                                                                                                                                                                                                                                       KLRSQLVKRK 586
                                                                                                                                                                                                                                                                                                                                                             Similarity
9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MANITOBA CANCER TREATMENT & RES FOUND. UNIV MANITOBA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       630 AA;
                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Turley
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279..382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       555..565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note=
                                                                 entry)
                                                                                                                                                                                                                                                                                                                                                                                  93.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              autoimmune disease; diagnosis; prognosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Hyaluronic acid binding domain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "repeat region"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      interaction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            631 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of cell-associated RHAMM with
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1; Mismatches
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motility), a protein involved in cell locomotion of moving cell proliferation and transformation. It differs from RHANM 1 by an insertion of 25 amino acids (see also AAR99674) between amino acids 54 and 55 of RHANM 1, resulting from an alternatively spliced exon 2A (AAT34502). RHANM 1-2a is the isoform that is overexpressed in tumours. Determination of the level of RHANM 1-2a in the insertion of the level of RHANM 1-2a in a sample can be used to assess the prognosis of a tumour less breast cancer) patient. The RHANM 1-2a protein can also (esp. breast cancer) patient. The RHANM 1-2a protein can also
                                                                    RHAMM; inflammatory neurological disorder; Parkinson's disease; althritis; multiple sclerosis; gastritis; nephritis; inflammatory dermatosis; psoriasis; inflammatory bowel disease;
                                                                                                                                                         Tissue disorder; response-to-injury process; cell proliferating; hyaluronic acid; HA; receptor for hyalauronan-mediated motility; RHAMM; inflammatory neurological disorder; Parkinson's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RHAMM 1-2a (AAR99675) is an alternatively spliced variant of RHAMM 1 (AAR99673) (receptor for hyaluronic acid mediated motility), a protein involved in cell locomotion or motility
                                  stenosis; restenosis; cancer; kidney fibrosis;
                                                                                                                                                                                                                                                                                                                       Mouse receptor for hyalauronan-mediated motility (RHAMM)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            a tumour by modulating the RHAMM with its ligand.
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                                                                                                                                                                                                                                                                                                                                                                                   CC hyalauronan-mediated motility (RHAMM), a polypeptide fragment encoding CC any of D1-D5 of RHAMM, or a vector which expresses antisense RHAMM, CC antibodies or a polypeptide fragment. The method is useful for treating a CC patient with an inflammatory neurological disorder such as Parkinson's CC disease, Alzheimer's disease, arthritis including rheumatoid arthritis, CC osteoarthritis, multiple sclerosis, inflammatory dermatosis (psoriasis), CC inflammatory bowel disease, stenosis or restenosis, cancer, kidney CC fibrosis, inflammatory lung disease (e.g. emphysema, asthma, cystic CC fibrosis, obesity or obesity related diseases, lupus, cardiovascular CC disease (e.g. atherosclerosis), and wound especially surgical excision CC disease (e.g. atherosclerosis), and wound especially surgical excision CC else skin grafte), stroke, inflammatory responses or fibrotic response CC associated with medical implants such as hip implants, vascular wraps and CC fibrosis, chronic cystitis, acute mastitis, gastritis, nephritis, CC hepatitis, septic shock, thyroiditis, and retinopathy. This sequence CC represents a receptor for hyalauronan-mediated motility protein used in the method of treating a tissue disorder described in the invention.
                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  with response-to-injury process or proliferating cells in a patient, comprising administering a polypeptide (I) which binds hyaluronic acid (HA), an antibody which binds one of domains D1-D5 of Receptor for (HA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treating tissue disorder associated with response-to-injury process or proliferating cells in mammals, e.g. fibrosis, inflammation, by administering a compound that alters activity of transition molecules
Mouse hyaluronic acid binding protein RHAMM
                                        12-MAR-2002
                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Fig 50; 215pp; English.
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                                                                             AAU11437;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 shock; thyroiditis; retinopathy.
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                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          describes a method of treating a tissue disorder associated
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                                                                                                                                                                                                                                                                                                93.5%;
                                                                                                                                                                                                                                                                                                  Score 43; 1
Pred. No. 5
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                                                                                                                                                                                                                                                                                            5.5;
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AIDS; hepatitis;
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The invention relates to a conjugate comprising an hyaluronic acid (HA) CC -binding protein e.g. RHAMM (receptor of HA mediated mobility) or peptide contiguous with, or coupled to a polypeptide conjugated to a therapeutic configuous with, or coupled to a polypeptide conjugated to a therapeutic configuration of the HA-binding protein by inserting a first nucleotide sequence encoding a HA-binding protein directly linked to a second conclectly expressing the vector in an acceptable host, purifying conjugate molecule from host or expression medium. The composition is useful for caltering in vivo the distribution of a therapeutic agent comprising conjugate endogenous HA and for tresting mammal where conjugate molecule will candinistering the composition to the animal where conjugate molecule will conjugate of the mammal contains high level of HA e.g. rheumatoid conjugates required also translates into lower immunogenicity of the conjugates improve patient compliance and reduce direct and indirect costs associated with the drug substance and reduce direct and indirect costs associated with the conventional dosage requirements for the conventional dosage requirements for the concasted balf-life and potency, resulting in prolonged circulation of the molecule, efficient distribution into the target tissues, and concressed balovavilability. The present seruence represents a RHAMM in the convent of the conventers of the structure of the structure of the conventers of the conventers of the structure of the conventers of the conventers of the conventers of the structure of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conventers of the conve
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Sequence
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                                                                                                                                                 CC This peptide represents a motif found in a binding domain of rat CC hyaluronan receptor corresponding to amino acid position 424-433. This creeptor is also known as the receptor for hyaluronic acid mediated motility (RHAMM). Hyaluronan is a large glycosaminoglycan that is call migration, growth and transformation. It interacts with call surfaces via specific protein receptors, e.g. RHAMM, that mediate many biological effects. The RHAMM/Hyaluronic acid interacts with call mological effects. The RHAMM/Hyaluronic acid interaction is involved in oncogene-and growth factor-mediated cell locomotion. The products can be used in the treatment of disorders, acute and chronic inflammatory disorders, Alzhelmer's and other forms of dementia, including Paskinson's and Huntington's diseases, AIDS, diabetes, autoimune diseases, corneal dysplasia and hypertrophies, burns, surgical including Paskinson's and Huntington's diseases, AIDS, diabetes, autoimune diseases, corneal dysplasia and hypertrophies, burns, surgical including Paskinson's and Huntington's diseases, AIDS, diabetes, autoimune diseases, corneal dysplasia and hypertrophies, burns, surgical inclusions and adhesions, strokes and multiple sclerosis. They can also be used in e.g. CNS and spinal cord regeneration, contraception and in vitro fertilitation and embryo development. The products can also be used in detection, diagnosis and prognosis.
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                                                                                             The invention relates to a conjugate comprising an hyaluronic acid (HA) C - binding protein e.g. RHAMM (receptor of HA mediated mobility) or peptide contiguous with, or coupled to a polypeptide conjugated to a therapeutic configuous with, or coupled to a polypeptide conjugated to a therapeutic configuous with, or coupled to a polypeptide conjugated to a therapeutic gener, and the polynucleotides encoding them. Also included is a method configure, and the polynucleotides encoding them. Also included is a method configure, and the polynucleotide sequence encoding protein directly linked to a second conclectle from host or expression medium. The composition is useful for altering in vivo the distribution of a therapeutic agent comprising conjugate molecule will distribute primarily in tissues and organs containing high levels of endogenous HA and for treating mammal where conjugate molecule will conjugate of the mammal contains high level of HA e.g. rheumatoid conjugated protein as compared to the native protein. As a result, conjugates improve patient compliance and reduce direct and indirect conjugates allows for the use, where appropriate, of lower, safer, conjugates allows for the use, where appropriate, of lower, safer, conjugates allows for the use, where appropriate, of lower, safer, conjugates and contency, resulting in prolonged circulation of the molecule, efficient distribution into the target tissues, and increased bioavailability. The present sequence represents a RHAMM increased bioavailability. The present sequence represents a RHAMM is a conjugate and potency, resulting in prolonged circulation of the molecule and potency.
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Query Match Best Local S Matches 8

Similarity 8; Conserv

Conservative

84.8%;

Score 39; DB Pred. No. 35; 1; Mismatches

DB 23; 35;

Length 713; Indels

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Gaps

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Sequence

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||:||| ||| 657 KLRSQLAKRK 666

1 KLKSQLVKRK 10

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Query Match
Best Local Similarity
Marches 8; Conserv?
                                                                                                                                     RESULT 15
AAR46550
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                                                                                                                                                                                The sequence is that of a binding motif fragment of the hyaluronan receptor (HARC). HARC is down regulated in normal cells and is only expressed in situations where cell motility is desired, e.g. in wound healing, in response to growth factors and in chemotaxis by white blood cells. HA may be used for diagnosis and treatment of diseases involving cell locomotion, e.g. tumour invasion, birth defects, acute and chronic inflammatory discorders, Alzheimer; s and other forms of dementia, ALDS, diabetes, autoimmure diseases, corneal dysplasias and hypertrophies, burns, surgical incisions and adhesions, strokes, multiple sclerosis, depression/schizophrenia related to neuronal growth and pain states involving nerve sprouting; also in CNJ and spinal cord regeneration, contraception, in vitro fertilisation and contraceptions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hyaluronan binding protein; HA; RHAMM; mediated motility; wound; healing; diagnosis; treatment; cell locomotion; tumour invasion; birth defects; inflammatory disorder; Alzheimer's disease; dementia; Parkinson's disease; Huntington's disease; AIDS; diabetes; auto; immune diseases; corneal dysplasia; hypertrophy; surgery; burns; strokes; multiple sclerosis; depression; schizophrenia; CNJ; contraception; in vitro fertilisation; embryo development.
                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 17; Page 57; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA encoding hyaluronan receptor - used to produce proteins and antibodies for alteration of cell locomotion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-APR-1992;
                                                                                                                                                                     See also AAR46548-51 and AAR43563.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1993-351722/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Turley EA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hyaluronan receptor binding motif (claimed).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-APR-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MANI-) MANITOBA CANCER TREATMENT & RES FOUND. (UYMA-) UNIV MANITOBA.
                                   1 KLKSQLVKR 9
KLRSQLVKR 9
                                                                                                                                     9 AA;
                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                  82.6%;
                                                               Score 38; DB 14; Length 9; Pred. No. 7.8e+05; 1; Mismatches 0; Indels
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Search completed: February 8, 2003, 10:41:01 Job time : 30 secs

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Result
No.
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Maximum
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Perfect score:
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                                                                                                                                                                                                                                                                                                                 score greater than or equal to the score of the result bein and is derived by analysis of the total score distribution.
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seq length: 2000000000
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Match
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- 31.670 Million cell updates/sec
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46
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Listing first 45 s
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                                                                                                                                                                                                                                                                                                                                                                 /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US60_PUBCOMB.pep:*
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                                        GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd
                                                                                                                                                                                                                                                        DB
                               US-10-012-896-592
US-09-895-793-592
US-09-895-814-592
US-09-895-814-592
US-09-875-143-592
US-09-759-143-592
US-09-864-761-33351
US-09-864-761-33351
US-09-864-761-33351
US-09-815-242-12463
US-09-815-242-12463
US-09-815-242-13094
US-09-815-242-131094
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US-09-815-242-131094
US-09-815-242-131094
US-09-1310-1318
US-10-029-009-11
US-110-029-009-11
US-10-029-009-11
US-10-029-110-63
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Sequence 11, Appl
Sequence 6, Appl
Sequence 26, Appli
Sequence 26, Appl
Sequence 1863, Ap
Sequence 5635, Ap
                                                                      Sequence 592, App
Sequence 592, App
Sequence 592, App
Sequence 592, App
Sequence 592, App
Sequence 593, App
Sequence 46169, A
Sequence 13351, A
Sequence 12433, A
Sequence 12667, A
Sequence 12094, A
Sequence 13094, A
Sequence 191, App
Sequence 191, App
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28	28	28	28	28	28	28	28	28	28	29	29	29	29	29	29	29	29	29	29	29	29	29	29	30	30
60.9	60.9	60.9	60.9	60.9	60.9	60.9	60.9	60.9	60.9	63.0	63.0	63.0	63.0	63.0	63.0	63.0	63.0	63.0	63.0	63.0	63.0	63.0	63.0	65.2	65.2
273	239	160	153	142	142	127	64	64	61	2165	2150	1185	909	901	835	621	603	498	478	460	460	421	229	3169	1224
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US-09-738-626-5061	US-09-764-898-165	US-09-764-864-1129	US-09-864-761-47313	US-09-738-973-212	US-09-854-133-212	US-09-764-868-801	US-09-815-242-11475	US-09-815-242-11296	US-09-864-761-34874	US-09-800-729-155	US-09-321-987B-2	US-09-895-913A-246	US-09-925-299-988	US-09-815-242-11148	US-09-801-574-22	US-09-856-247A-2	US-09-864-761-36676	US-10-029-009-22	US-10-029-009-10	US-09-884-430-6	US-09-782-980-82	US-09-841-132-577	US-09-764-898-253	US-10-114-170-257	US-09-801-368-222
5061,	165,	Sequence 1129, Ap	\vdash	e 21		108				15	æ	246	Sequence 988, App		N		e 3	Sequence 22, Appl		Sequence 6, Appli	82	Sequence 577, App	Sequence 253, App	Sequence 257, App	Sequence 222, App

ALIGNMENTS

RESULT 1

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APPLICANT: Meagher, Madeleine Joy
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE REFERENCE: 210121.427C27
CURRENT APPLICATION NUMBER: US/10/012,896
CURRENT FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 1011
SOPTWARE: FRSTSEQ for Windows Version 3.0
SEQ ID NO 592
LENGTH: 188
TYPE: PRT
ORGANISM: Homo sapiens
US-10-012-896-592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-012-896-592; Sequence 592; Application No. US20; Publication No. US20; GENERAL INFORMATION:
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
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APPLICANT:
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Vedvick, Thomas
Carter, Darrick
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McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
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Hepler, William T.
Henderson, Robert A.
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Kalos, Michael D.
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CURRENT APPLICATION NUMBER: US/09/895,793
CURRENT FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FRSEUSEQ for Windows Version 3.0
SEQ ID NO 592
LENGTH: 188
TYPE: PRT
ORGANISM: Homo sapien
US-09-895-793-592
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Best Local Similarity
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                                                                                                                                               Sequence 592, Application US/09895814 Publication No. US20020193296A1 GENERAL INFORMATION:
                                             APPLICANT: Xu, CAPPLICANT: Dill APPLICANT: Harl APPLICANT: Harl APPLICANT: Kall APPLICANT: Kall APPLICANT: Rett
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Best Local Similarity
Matches 7; Conserv
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.534C2
                  PPLICANT:
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                                                                                                                                                                                                                                                       178 LKAQLVKQK 186
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                                                                                                                                                                                                                                                                                  2 LKSQLVKRK 10
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             Jiang, Yugiu
Kalos, Michael D.
Retter, Marc W.
Stolk, John A.
Day, Craig H.
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Kalos, Michael D.
Retter, Marc W.
Stolk, John A.
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Carter, Darrick
Li, Samuel X.
                                                                                       Mitcham, Jennifer L.
Harlocker, Susan L.
Day, Craig H.
Vedvick, Thomas S.
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Hepler, William T.
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Vinals de Bassols, Carlota
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o. US20020192763A1
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77.8%;
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; Pred. No. 16;
2; Mismatches
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Pred. No.
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18
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                                                                                                                                                                                                                                                                                                                                         Length 188;
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Query Match
Best Local Similarity
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                                                                            ; LENGTH: 188
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-759-143-592
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US-09-895-814-592
                                                                                                                                     APPLICANT: Skelky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIACNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C23
CURRENT APPLICATION NUMBER: US/09/759,143
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 592
SEQ ID NO 592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/895,814
CURRENT FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 990
SOPTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 592
LENGTH: 188
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 592, Application US/09759143
Patent No. US20020022248A1
GENERAL INFORMATION:
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Best Local Similarity 77.8%;
Matches 7; Conservative
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APPLICANT:
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APPLICANT:
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C26
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Vedvick, Thomas S
Carter, Darrick
Li, Samuel
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Hepler, William T.
Henderson, Robert A.
Hural, John
McNeill, Patricia D.
Houghcon, Raymond L.
Vinals de Bassols, Carlota
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Mitcham, Jennifer
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              Conservative
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Pred. No. 16;
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APPLICANT: HOUGHTON, KAYMOND AND METHODS
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CI-
FILE REFERENCE: 210121.427C24
CURRENT APPLICATION NUMBER: US/09/780,669
CURRENT FILING DATE: 2001-02-09
NUMBER OF SEQ ID NOS: 943
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 592
LENGTH: 188
TWORD: DRT
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US-09-780-669-592
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Matches
                                                                                         SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 592
LENGIH: 188
                                                                                                                                                                                                                                                Sequence 592, Application Patent No. US20020081680A1 GENERAL INFORMATION:
 Best Local Similarity
                 Query Match
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                                                                                                                                     APPLICANT: XI, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.534C1
CURRENT FILING DATE: 2001-03-28
CURRENT FILING DATE: 2001-03-28
UNMBER OF SEQ ID NOS: 982
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                                               ORGANISM: Homo sapien
-09-822-827-592
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Houghton, Raymond L.
FANTION: COMPOSITIONS AND METHODS FOR
FENTION: DIAGNOSIS OF PROSTATE CANCER
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Hepler, William
Hural, John
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Kalos, Michael D.
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Mitcham, Jennifer L.
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Carter, Darrick
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SOFTWARE: Annomax Sequence Listing
SEQ ID NO 46169
LENGTH: 55
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CURRENT FILING DATE: 2001-05-23
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PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
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PRIOR APPLICATION NUMBER: PCT/US01/00666
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PRIOR APPLICATION NUMBER: US 60/236,359
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PRIOR APPLICATION NUMBER: GB 24263.6
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PRIOR FILING DATE: 2000-05-26
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PRIOR APPLICATION NUMBER: PCT/US01/00670
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OTHER INFORMATION: MAP TO ACO09275.5
OTHER INFORMATION: EXPRESSED IN LING, SIGNAL = 3.3
OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 2.7
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.4
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 2.9
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 2.9
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 2.9
OTHER INFORMATION: EXT_HUMAN HIT: BE257210.1, EVALUE 4.00e-25
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                                                                                                                                                                        ORGANISM: Homo sapiens FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: PCT/US01/00661
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Hanzel, David K.
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PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 99/632,366
PRIOR APPLICATION NUMBER: US 99/632,366
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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                                                            PEATURE:
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IEQ ID NO 33351

IEQNITH: 69

TYPE: PRT
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CURRENT FILING DATE: 2001-05-23
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                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                  OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00670
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: US 60/234,687
OR FILING DATE: 2000-09-21
OR APPLICATION NUMBER: US 09/608,408
OR FILING DATE: 2000-06-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OR APPLICATION NUMBER: US 60/236,359
OR FILING DATE: 2000-09-27
OR APPLICATION NUMBER: PCT/US01/00666
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00667
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00664
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les 6; Conserv
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                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 09/774,203 FILING DATE: 2001-01-29
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APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: PCT/US01/00661
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/US01/00662 FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: PCT/US01/00663 FILING DATE: 2001-01-30
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5. US20020048763A1
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Hanzel, David K.
Chen, Wensheng
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N: MAP TO AC006160.9

N: EXPRESSED IN BT474, SIGNAL = 1.1

N: EXPRESSED IN HELA, SIGNAL = 1.2

N: EXPRESSED IN BRAIN, SIGNAL = 1.3

N: EXPRESSED IN PLACENTA, SIGNAL = 0.6

N: EXPRESSED IN HEL100, SIGNAL = 1.1

N: EXPRESSED IN FETAL LIVER, SIGNAL =
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Pred. No.
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Sequence 12867, Application US/
Patent No. US20020061569A1
GENERAL INFORMATION:
APPLICANT: Haselbeck, Robert
APPLICANT: Ohleen, Kari L.
APPLICANT: Zyskind, Judith W.
                                                                                                                                                                                                                        S
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                                                                                                             US-09-815-242-12867
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CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
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SOFTWARE: FastSEQ for
SEQ ID NO 12433
LENGTH: 117
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PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 60/269,308
PRIOR FILING DATE: 2001-02-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
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                                                                                                                                                                                                                                                                                  Local
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7; Conserva
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7; Conserv
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Zyskind, Judith
Wall, Daniel
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IN ADULT LIVER, SIGNAL = 1
IN HEART, SIGNAL = 0.82
HIT: BE168080.1, EVALUE 4.00e-04
HIT: P25386, EVALUE 2.00e-03
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Pred. No.
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Pred. No. 13;
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APPLICANT: Haselbeck, Robert
APPLICANT: Ohleen, Kari L.
APPLICANT: Ohleen, Kari L.
APPLICANT: Ohleen, Kari L.
APPLICANT: Zyskind, Judith W.
APPLICANT: Wall, Daniel
APPLICANT: Trawick, John D.
APPLICANT: Trawick, John D.
APPLICANT: Yamamoto, Robert T.
APPLICANT: Yamamoto, Robert T.
APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT APPLICATION NUMBER: 60/191,078
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
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US-09-815-242-12867
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CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
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Best Local Similarity
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                                                PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
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PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
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NITLE OF INVENTION: Identification of Essential Genes
NITLE OF INVENTION: Prokaryotes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
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FILING DATE: 2000-12-22
APPLICATION NUMBER: 60/269,308
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70.0%;
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Pred. No.
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RESULT 13
US-09-729-674-158
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; ORGANISM: Homo sapiens
US-09-789-561-191
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CURRENT FILING DATE: 2001-02-22
PRIOR APPLICATION NUMBER: PCT/US00/24008
PRIOR FILING DATE: 2000-08-31
PRIOR APPLICATION NUMBER: 60/152,317
PRIOR APPLICATION NUMBER: 60/152,315
PRIOR FILING DATE: 1999-09-03
PRIOR FILING DATE: 1999-09-03
PRIOR FILING DATE: 1999-09-03
PRIOR FILING DATE: 1999-09-03
PRIOR FILING DATE: 1999-09-03
                                                                                                                                                                                                                                                                                          Sequence 158, Application US/09729674
Patent No. US20010039335A1
GENERAL INFORMATION:
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SOPTWARE: FRSTSEQ for Windows Version 4.0
SEQ ID NO 13094
LENGTH: 117
TYPE: PRT
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Best Local Similarity
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APPLICANT:
APPLICANT:
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CURRENT APPLICATION NUMBER: US/09/729,674
CURRENT FILING DATE: 2000-12-04
                              APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING
FILE REFERENCE: 6055-64X
                                                                                                                                                                                                                                                           APPLICANT: Jacobs, Kenneth APPLICANT: McCoy, John M.
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FILE REFERENCE: PZO
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                                                                             Clark, Hilary
Fechtel, Kim
                                                                                                          Steininger II, Robert
Spaulding, Vikki
Wong, Gordon G.
                                                                                                                                                             Agostino, Michael
                                                                                                                                                                          Merberg, David
Treacy, Maurice
                                                                                                                                                                                                              Evans, Chery:
                                                                                                                                                                                                                           Collins-Racie, Lisa
                                                                                                                                                                                                                                            LaVallie, Edward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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Pred. No.
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Pred. No. 28;
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US-10-029-009-23

US-10-029-009-23

Sequence 23, Application US/10029009

Patent No. US20020164617A1

GENERAL INFORMATION:

APPLICANT: Felsch, Jason S.

APPLICANT: Kalghatgi, Krishna

APPLICANT: Nash, Huw M.

TITLE OF INVENTION: Affinity Selection-Based Screening of Hydrophobic Proteins

FILE REFERENCE: 111025-173 US2

CURRENT APPLICATION NUMBER: US/10/029,009

CURRENT FILING DATE: 2002-03-28

PRIOR APPLICATION NUMBER: US 60/258,970

PRIOR APPLICATION STATE: 2006-12-29

NUMBER OF SEO ID NOS: 34

SÖFTWARE: FastSEQ for Windows Version 4.0

SEQ ID. NO 23

LENGTH: 626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Affinity Selection-Based Screening of Hydrophobic Proteins
FILE REFERENCE: 111025.173 US2
CURRENT APPLICATION NUMBER: US/10/029,009
CURRENT FILING DATE: 2002-03-28
PRIOR APPLICATION NUMBER: US 60/258,970
PRIOR FILING DATE: 2000-12-29
NUMBER OF SEQ ID NOS: 34
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11
LENGTH: 608
TYPE: PRT
ORGANISM: Rat
US-10-029-009-11
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US-10-029-009-11
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Best Local Similarity 60.0%;
Matches 6; Conservative
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Best Local Similarity 60.0%;
Matches 6; Conservative
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PRIOR FILING DATE: 2000-03-30
NUMBER OF SEQ ID NOS: 283
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 158
LENGTH: 586
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Felsch, Jason S.
APPLICANT: Annis, David Allen
APPLICANT: Kalghatgi, Krishna
APPLICANT: Nash, Huw M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: UNSURE
LOCATION: (286)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  469 KTRSQITKRK 478
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Pred. No. 1.2e+02;
3; Mismatches 1
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; TYPE: PRT
; CORGANISM: Rat
US-10-029-009-23

Query Match
Best Local Similarity 60.0%; Pred. No. 1.3e+02;
Matches 6; Conservative 2; Mismatches 2; Indels 0; Gaps

Qy 1 KLKSQLVKRK 10
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Db 485 KTRSQITKRK 494
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Search completed: February Job time : 9 secs

8, 2003, 10:52:11

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Minimum
Maximum
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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seq length: 2000000000
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48.476 Million cell updates/sec
                 4569144 seqs, 644733110 residues
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/ Cgm2_6/ptodata/1/paa/US06_COMB.pep: *
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/ Cgm2_6/ptodata/1 paa/US06_COMB.pep: *
/ Cgm2_6/ptodata/1 paa/US080_COMB.pep: *
/ Cgm2_6/ptodata/1 paa/US081_COMB.pep: *
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/ Cgm2_6/ptodata/1 paa/US084_COMB.pep: *
/ Cgm2_6/ptodata/1 paa/US086_COMB.pep: *
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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44444 00000	
100.0 100.0 100.0 100.0 93.5 1 93.5 1	Query Match
100	Query Match Length DB ID
7 220 15 18	DB.
7 US-08-318-892-4 20 US-09-685-010-31 23 US-09-978-309A-31 7 US-08-318-892B-4 15 US-09-169-077-2 18 US-09-424-292-6	ID
Sequence 4, Appli Sequence 31, Appl Sequence 31, Appl Sequence 4, Appli Sequence 2, Appli Sequence 6, Appli	Description

45	44	43	42	41	40	39	38	37	36	35	34	မ	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	80	7
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US-09-563-582-54	44-	-10-155-88	9-816-660-	09-733-089-	-09-424-292-	2-3	-2	318-892B-1	18-892-13	-978	ů	8-318-892B-	-10-023-634-8	-09-169-077	-09-791-	US-09-978-309A-48	685-010-	09-169-	09-791-537-673	US-09-791-537-88053	US-08-318-892B-2	US-09-978-309A-79	09-978-309A-7	US-09-791-537-42360	8-318-8	-309A	8-318-8		-09-978-	-09-978-	-09-978-309A-	-09-685-010-5	-09-978-	-010-2	08-318-8		-09-424-292-2	9
Sequence 54, Appl	Sequence 8435, Ap	33553,	8991,	8991,	32, A	30,	e 23,	13, 7	13, /	59,	e 59,	32	18	~	22	48,	48,		673	е 8	N)	7	7	e 42	.~	e 80,	33,	73,	75,	e 82,	e 51,	e 51,	e 24,	e 24	10,		e 28,	-

ALIGNMENTS

US-08-318-892-4

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Sequence 4, Application US/08318892
GENERAL INFORMATION:
APPLICANT: TULLEY, EVA A
TITLE OF INVENTION: HYALURONAN RECEPTOR (RHAMM = RECEPTOR
TITLE OF INVENTION: POR HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING
TITLE OF INVENTION: PEPTIDES
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSE: c/o FISH & NEAVE
STREET: 1251 Avenue of the Americas
CITY: New York
STATE: N.Y.
COUNTRY: USA
ZIP: 10020
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/318,892
FILING DATE: 1-DEC-1994
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Pierri, Margaret A
REGISTRATION NUMBER: 30,709
REFERENCE/DOCKET NUMBER: SIM-8
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APPLICANT: Cruz, Tony
APPLICANT: Cruz, Tony
APPLICANT: Pastrak, Aleksandra
APPLICANT: Turley, Eva A.
TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response to
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated by
TITLE OF INVENTION: Hyaladherin and Hyaluronans
FILE REFERENCE: 033352-010
CURRENT APPLICATION NUMBER: US/99/978,309A
CURRENT FILING DATE: 2002-04-04
PRIOR APPLICATION NUMBER: US 09/685,010
PRIOR APPLICATION NUMBER: US 09/541,522
PRIOR APPLICATION NUMBER: US 09/541,522
PRIOR APPLICATION NUMBER: US 09/541,522
PRIOR APPLICATION NUMBER: US 09/541,522
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TITLE OF INVENTION: CELLULAR RESPONSE TO INJURY AND OTHER PROLIFERATING CELL
TITLE OF INVENTION: DESCRIPTIONS TO INJURY AND OTHER PROLIFERATING CELL
TITLE OF INVENTION: CELLULAR RESPONSE TO INJURY AND OTHER PROLIFERATION CELL
TITLE OF INVENTION: DIVERSE PROLIFERATION NUMBER REGULATED BY HYALADHERIN AND HYALJURONANS
FILE REFERENCE: 9.0130,401C1
CURRENT PILLING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 72
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 31
LEUGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 31, Application US/09978309A GENERAL INFORMATION:
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Best Local Similarity
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LENGTH: 10 amino acids
TYPE: amino acid
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STRANDEDNESS: Bin
STRANDEDNESS: Bin
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les 10; Conserv
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SEQ ID NOS: 84
FASTSEQ for Windows Version 4.0
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100.0%; Pr
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Pred. No. 0.11;
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Pred. No. 0.11;
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US-09-169-077-2
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; OTHER INFORMATION: Peptide composition that binds a hyalauronan US-09-978-309A-31
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                                                                                                                                              Best Local Similarity Matches 9; Conserv
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GENERAL INFORMATION:
APPLICANT: Turley
Sequence 2, Application US/09169077 GENERAL INFORMATION:
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LENGTH: 10
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Best Local Similarity
Matches 10; Conserv
                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/318
PILING DATE: 1-DEC-1994
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Pierri, Margaret A
REGISTRATION NUMBER: 30709
REFERENCE/DOCKET NUMBER: SIM-
TELECOMMUNICATION INFORMATION:
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: FORM PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                         MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                  TYPE: amino o
                                                                                                                                                                                                                                          TOPOLOGY:
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amino acid
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1251 Avenue of the Americas
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Pred. No.
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Pred. No.
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TITLE OF INVENTION: NUMBER OF SEQUENCES:

HUMAN HYALURONAN RECEPTOR 52

APPLICANT:

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Query Match
Best Local Similarity
"hes 9; Conserv?
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US-09-424-292-6
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                                                                                                                         US-09-424-292-15
                                                                                                                                           RESULT 7
                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Synthetic US-09-424-292-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6, Application US/09424292
GENERAL INFORMATION:
APPLICANT: Turley, Eva A
TITLE OF INVENTION: Improved Delivery of Disease Modifiers
FILE REFERENCE: 205375
CURRENT APPLICATION NUMBER: US/09/424,292
CURRENT FILING DATE: 2000-07-06
                                                                                      Sequence 15, Application US/09424292 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 6
LENGTH: 10
                                                                                                                                                                                                                                                                    Matches
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APPLICANT: Turley, Eva A
TITLE OF INVENTION: Improved Delivery of Disease Modifiers
FILE REFERENCE: 205375
CURRENT APPLICATION NUMBER: US/09/424,292
CURRENT FILING DATE: 2000-07-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: CA 2,205,771
PRIOR FILING DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 2000-07-06
PRIOR APPLICATION NUMBER: PCT/CA98/00448
PRIOR FILING DATE: 1998-05-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: GB 9607441.4
PILING DATE: 10-APR-1996
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/169,077
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/CA97/0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                  Local Similarity 90.
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TOPOLOGY: linear
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Pred. No. 0.42;
1; Mismatches
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Pred. No. 0.42;
1; Mismatches
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CURRENT APPLICATION NUMBER: US/09/424,292
CURRENT FILING DATE: 2000-07-06
PRIOR APPLICATION NUMBER: PCT/CA98/00448
PRIOR FILING DATE: 1998-05-11
PRIOR APPLICATION NUMBER: CA 2,205,771
PRIOR APPLICATION NUMBER: CA 2,205,771
PRIOR FILING DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 36
SOPTWARE: Patentin version 3.1
SEQ ID NO 28
                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 9; Conserva
                                                                                                                                                 SEQUENCE 10, Application US
GENERAL INFORMATION:
APPLICANT: Turley, Eva.
TITLE OF INVENTION: HYA.
TITLE OF INVENTION: FOR
TITLE OF INVENTION: POP
TITLE OF INVENTION: PEP
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSES
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SOFTWARE: Patentin version 3.1
SEQ ID NO 15
LENGTH: 10
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Matches 9; Conserv
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PRIOR APPLICATION NUMBER: CA 2,205,771
PRIOR FILING DATE: 1997-05-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Turley, Eva A
TITLE OF INVENTION: Improved Delivery of Disease Modifiers
FILE REFERENCE: 205375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
OTHER INFORMATION: Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                      CITY: New York STATE: N.Y.
                                                    COUNTRY: U
                                                                                                                        STREET:
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1251 Avenue of the Americas
                                                                       USA
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    Mismatches

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Pred. No. 0.
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Pred. No. 0.42;
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RESULT 10
US-08-318-892B-10
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Best Local Similarity
Finds 9; Conserve
                                                                 US-08-318-892B-10
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                                                                                                                                                             CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Pietri, Margaret A
REGISTRATION NUMBER: 30709
REFERENCE/DOCKET NUMBER: SIM-
TELECOMMUNICATION INFORMATION:
TELEPAN: (212) 596-9000
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 10, Application US/08318892B
GENERAL INFORMATION:
APPLICANT: Turley, Eva A
TITLE OF INVENTION: HYALURONAN REC
TITLE OF INVENTION: HYALURONAN MEI
NUMBER OF SEQUENCES: 41
Best Local Similarity
                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (212) 596-9090
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                              STRANDEDNESS:
TOPOLOGY: line
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Pierri, Margaret A
REGISTRATION NUMBER: 30,709
REFERENCE/DOCKET NUMBER: SIM-8
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/01 FILING DATE: 1-DEC-1994 CLASSIFICATION: 536 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
                                                                                                                                   TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 1-DEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U
ZIP: 10020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: N.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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1251 Avenue of the Americas
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                                                                                                    linear
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                                                                                  peptide
                                                                                                                 single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HYALURONAN RECEPTOR (RHAMM = RECEPTOR FOR HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING PEPTIDES : 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93.5%;
            93.5%;
90.0%;
                                                                                                                                                                                                                                                                                                                                                                 US/08/318,892B
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                                                                                                                                                                                                                                                              8-MIS
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            Score 43; DB 7;
Pred. No. 0.47;
                           Length 11;
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APPLICANT: Tony
APPLICANT: Turley, Eva A.
APPLICANT: Turley, Eva A.
APPLICANT: Turley, Eva A.
APPLICANT: Turley, Eva A.
APPLICANT: Turley, Eva A.
TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response to
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated by
TITLE OF INVENTION: Hyaladherin and Hyaluronans
FILE REFERENCE: 033352-010
CURRENT APPLICATION NUMBER: US/09/978,309A
CURRENT FILING DATE: 2002-04-04
PRIOR APPLICATION NUMBER: US 09/685,010
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 2000-04-03
PRIOR APPLICATION NUMBER: US 60/127,457
PRIOR APPLICATION NUMBER: US 60/127,457
PRIOR APPLICATION NUMBER: US 60/127,457
PRIOR APPLICATION DATE: 199-04-01
NUMBER: OF SEO ID NOS: 84
SEC ID NO 24
LENGTH: 11
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US-09-685-010-24
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APPLICANT: Turley, Eva A.

APPLICANT: Cruz, Tony F.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATING
TITLE OF INVENTION: CELLULAR RESPONSE TO INJURY AND OTHER PROLIFERATING CELL
TITLE OF INVENTION: DISORDERS REGULATED BY HYALADHERIN AND HYALURONANS
FILE REFERENCE: 910130.401C1
CURRENT APPLICATION NUMBER: US/09/685,010
CURRENT APPLICATION NUMBER: US/09/685,010
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 72
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 24
LENGTH: 11
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Best Local Similarity 90.0
Thes 9; Conservative
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                                                                            Matches
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                                                                          Best Local Similarity
Matches 9; Conserv
                                                                                                               Query Match
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ORGANISM: Homo sapien
                                                                                                                                                                         ORGANISM: Homo sapien
                                    1 KLKSQLVKRK 10
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1 KLRSQLVKRK 10
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                                                                          Conservative
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                                                                                            93.5%;
                                                                        Score 43; DB 2
Pred. No. 0.47;
1; Mismatches
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Pred. No. 0.
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RESULT 13

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RESULT 15
US-09-978-309A-82
; Sequence 82, Application US/09978309A
; GENERAL INFORMATION:
; APPLICANT: Cruz, Tony
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; OTHER INFORMATION: Peptide used in competition binding assay
US-09-978-309A-51
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CURRENT APPLICATION NUMBER: US/09/978,309A
CURRENT FILING DATE: 2002-04-04
PRIOR APPLICATION NUMBER: US 09/685,010
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 2000-00-05
PRIOR APPLICATION NUMBER: US 09/541,522
PRIOR FILING DATE: 2000-04-03
PRIOR APPLICATION NUMBER: US 60/127,457
PRIOR FILING DATE: 1999-04-01
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Best Local Similarity 90.0
Watches 9; Conservative
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GENERAL INFORMATION:
APPLICANT: Cruz, Tony
APPLICANT: Pastrak, Aleksandra
APPLICANT: Turley, Eva A.
TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response to
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated by
TITLE OF INVENTION: Hyaladherin and Hyaluronans
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Matches
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APPLICANT: Cruz, Tony F.
APPLICANT: Cruz, Tony F.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATING
TITLE OF INVENTION: CELLULAR RESPONSE TO INJURY AND OTHER PROLIFERATING CELL
TITLE OF INVENTION: DISORDERS REGULATED BY HYALADHERIN AND HYALURONANS
FILE REFERENCE: 910130.401C1
CURRENT APPLICATION NUMBER: US/09/685,010
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 72
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 51
LENGTH: 32
TYPE: THE TIME OF THE TOTAL PROPERTY OF THE TOTAL PROPERTY OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME OF THE TIME
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 51
LENGTH: 32
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ORGANISM: Unknown
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23 KLRSQLVKRK 32
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Pred. No. 1.6;
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Search completed: February Job time: 133 secs
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CURRENT APPLICATION NUMBER: US/09/978,309A

CURRENT FILING DATE: 2002-04-04

PRIOR APPLICATION NUMBER: US 09/685,010

PRIOR FILING DATE: 2000-10-05

PRIOR PILING DATE: 2000-04-03

PRIOR PILING DATE: 2000-04-03

PRIOR PILING DATE: 2000-04-03

PRIOR APPLICATION NUMBER: US 60/127,457

PRIOR PILING DATE: 1999-04-01
                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 84
SOFTWARE: FeatSEQ for Windows Version 4.0
SEQ ID NO 82
LENGTH: 32
                                                                                                                                                               Best Local Similarity Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Pastrak, Aleksandra
APPLICANT: Turley, Eva A.
TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response to
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated by
TITLE OF INVENTION: Hyaladherin and Hyaluronans
                                                                                                                                                                                                                                                                                                     TYPE: PRT
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                                                                                                                                                                      Conservative
                                                                                                                                                                                          93.5%;
90.0%;
                   8, 2003, 10:50:49
                                                                                                                                                                                          Score 43;
Pred. No.
                                                                                                                                                                        Mismatches
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Result
No.
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Maximum
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seq length: 2000000000
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   %
Query
Match
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1: /cgn2_6/ptodata/2/paa/PCT_NEW_COMB.pep:*
2: /cgn2_6/ptodata/2/paa/USO6_NEW_COMB.pep:*
3: /cgn2_6/ptodata/2/paa/USO7_NEW_COMB.pep:*
4: /cgn2_6/ptodata/2/paa/USO9_NEW_COMB.pep:*
5: /cgn2_6/ptodata/2/paa/USO9_NEW_COMB.pep:*
6: /cgn2_6/ptodata/2/paa/USO9_NEW_COMB.pep:*
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                               US-10-310-154-439
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29	29	29	29	29	29	29	29	29	29	29	30	30	30	30	30	30	30	30
63.0	63.0	63.0	63.0	63.0	63.0	63.0	63.0	63.0	63.0	63.0	65.2	65.2	65.2	65.2	65.2	65.2	65.2	65.2
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Sequence 61733, A	Sequence 61734, A	Sequence 61733, A	Sequence 6446, Ap	Sequence 639, App	Sequence 635, App	Sequence 6648, Ap	Sequence 6648, Ap	Sequence 3928, Ap	Sequence 11254, A	Sequence 11254, A	Sequence 4056, Ap	Sequence 2042, Ap	Sequence 381, App	Sequence 5248, Ap	Sequence 80, Appl	Sequence 4086, Ap	Sequence 4232, Ap	Sequence 4232, Ap

ALIGNMENTS

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RESULT 1
US-10-310-154-439
Sequence 439, Application US/10310154
GENERAL INFORMATION:
APPLICANT: Edgerton, Michael D
APPLICANT: Chomet, Paul S.
APPLICANT: Adams, Thomas H
APPLICANT: Ruff, Thomas G.
APPLICANT: Agarwal, Ameeta K.
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Luthy, Michael M.
Lund, Adrian
Madson, Linda L.
Malloy, Kathleen A.
McKiel, Christine L.
Miller, Phillp W.
Padmayathi, Manchikanti
                                                                                                                                                                                                                                                                                                                                                          Duff, Stephen M.
Galligan, Meghan M.
Hinchey, Brenda S.
Huang, Shihshieh
           Wang, Haiyun
Xin, Zhanguo
Xu, Nanfei
Yang, Chunzhi
Zeng, Xiaoping
Zhang, Qiang
Zhao, Yajuan
                                                                                                  Vidya,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bell,
                                                                                                   Tennesen, υ
Vidya, K.R.
                                                                                                                           Parnell, Laurence D. Start, William G.
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Laccetti, Lucille
Lai, Chao-Qiang
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Deikman, Jill
Deng, Molian
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Ahrens, Jeffrey
Ball, James A.
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; TITLE OF INVENTION: Gene Sequences and Uses Thereof in Plants
; FILE REPERENCE: 38-15(52796)B
; CURRENT APPLICATION NUMBER: US/10/310,154
; CURRENT FILING DATE: 2002-12-04
; PRIOR APPLICATION NUMBER: 60/337,358
PRIOR PILING DATE: 2001-12-04
; NUMBER OF SEQ ID NOS: 736
; SEQ ID NO 439
; SEQ ID NO 439
; SEQ ID NO 439
; TYPE: PRT
; ORGANISM: Zea mays
US-10-310-154-439
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US-60-434-832-7073
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SEQ ID NO 7073
LENGTH: 544
TYPE: PRT
ORGANISM: Candida albicans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1700,
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                    APPLICANT: SUWA, MAKIKO
APPLICANT: ASAI, KIYOSHI
APPLICANT: AKIYAMA, YUTAKA
APPLICANT: AKIYAMA, YUTAKA
APPLICANT: ABURATANI, HIROYUKI
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 084335/0152
CURRENT APPLICATION NUMBER: US/10/017,161
CURRENT FILLING DATE: 2002-12-18
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PRIOR FILING DATE: 2001-06-18
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APPLICANT: Jiang, Bo
APPLICANT: Boone, Charles
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                       TYPE: PRT
ORGANISM: Homo sapiens
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Pred. No. 11;
0; Mismatches
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US-10-294-025-592
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APPLICANT: Ensign, Jerald C
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 592
LENGTH: 188
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GENERAL INFORMATION:
GENERAL INFORMATION:
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GENERAL INFORMATION:
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GENERAL INFORMATION:
APPLICANT: Xu Jiangchun A.
APPLICANT: Xalos, Michael D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C29
CURRENT APPLICATION NUMBER: US/10/294,025
CURRENT FILLING DATE: 2002-11-12
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Best Local Similarity
Matches 7; Conserv
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                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/284,060
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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TITLE OF INVENTION: Insecticidal Protein Toxins from
TITLE OF INVENTION: Photorhabdus
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
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CITY: Indianapolis
STATE: IN
                                FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                       COUNTRY: US
ZIP: 46268
                                                                                                                                                                                                                                                                          ADDRESSEE:
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9330 Zionsville Road
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Merlo, Donald J
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Fatig, Raymond
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ffrench-Constant, Richard
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Bowen, David J
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77.8%;
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 US 08/063,615
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Pred. No.
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Pred. No. 5.
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FILING DATE:

18-MAY-1993

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US-10-284-060-28

: Sequence 28, Application US/10284060

: GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/284,060
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 50
TELECOMMUNICATION INFORMATION:
TELEPHONE: 317-337-4846
                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                TITLE OF INVENTION: Insecticidal TITLE OF INVENTION: Photorhabdus NUMBER OF SEQUENCES: 88
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APPLICATION NUMBER: US 06
FILING DATE: 28-AUG-1996
PRIOR APPLICATION DATA:
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ATTORNEY/AGENT INFORMATION:
NAME: BOTUCKI, Andrea T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/608,423
FILING DATE: 28-FEB-1996
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146 KLKSELVESK 155
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APPLICATION NUMBER:
FILING DATE: 28-FEB-
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Local Similarity 70.0%;
hes 7; Conservative
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                                                                                                                                                                                  STATE:
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CLASSIFICATION:
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9330 Zionsville Road
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Roberts, Jean I
Rocheleau, Thomas A
VENTION: Insecticidal Protein Toxins from
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Bowen, David J
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Fatig, Raymond
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06-NOV-1995
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Pred. No. 57
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; LENGTH: 627 amino acids
; TYPE; amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-10-284-060-28
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            ZIP: 46268

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
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                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: DowElanco
STREET: 9330 Zionsville Road
                                                                                                                                                                                                                                               APPLICANT: Rocheleau, Thomas A TITLE OF INVENTION: Insecticidal TITLE OF INVENTION: Photorhabdus
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LENGTH: 627 amino acid
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APPLICATION NUMBER:
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PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
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CITY: Indianapolis
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REGISTRATION NUMBER: 33:
REFERENCE/DOCKET NUMBER:
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APPLICATION NUMBER:
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                                                                                                                                COUNTRY:
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Rocheleau, Thomas
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Fatig, Raymond
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Bowen, David J
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70.0%;
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US/10/284,060
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 33; DB 6;
Pred. No. 57;
2; Mismatches
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SEQUENCE CHARACTERISTICS:
LENGTH: 627 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 26, Application US/10284060 GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
APPLICANT:
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TITLE OF IN
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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APPLICANT:
APPLICANT:
APPLICANT:
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ADDRESSEE: DowElanco
                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Borucki, Andrea T.
REGISTRATION NUMBER: 33651
REFERENCE/DOCKET NUMBER: 50301E
TELECOMMUNICATION INFORMATION:
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FILING DATE: 28-FEB-
PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/395,497
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ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
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PRIOR APPLICATION NUMBER: US 08/063,615
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                                                                                                                                   CITY: Indianapolis
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T: Rocheleau, Thomas A
INVENTION: Insecticidal Protein Toxins from
INVENTION: Photorhabdus
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Merlo, Donald J
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Bowen, David J
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28-FEB-1996
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; Pred. No. 57;
2; Mismatches
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; Sequence 26, Application US/10284060
; GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
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APPLICATION NUMBER: US 08/395,497
FILING DATE: 28-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/007,255
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                    TITLE OF INVENTION: Insecticidal TITLE OF INVENTION: Photorhabdus NUMBER OF SEQUENCES: 88 CORRESPONDENCE ADDRESS:
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APPLICANT:
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APPLICATION NUMBER: US 08/063,615
PILING DATE: 18-MAY-1993
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FILING DATE: 28-FEB-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/705,484
FILING DATE: 28-AUG-1996
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APPLICATION NUMBER: US/10/284,060
FILING DATE:
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APPLICATION NUMBER: US 08/743,699
FILING DATE: 06-NOV-1996
ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
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                                                     ZIP: 46268
                                                                                                  STREET: 9330 Zions CITY: Indianapolis
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                                                                   COUNTRY:
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es 7; Conserv
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                                                                                                                                      ADDRESSEE:
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REGISTRATION NUMBER: 33
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                                                                                                                   9330 Zionsville Road
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Rocheleau, Thomas A
VENTION: Insecticidal Protein Toxins
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Fatig, Raymond
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Bowen, David J
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Pred. No. 1.2e+0
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SOPTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:

US/10/284,060

OPERATING SYSTEM: PC-DOS/MS-DOS

FILING DATE: APPLICATION NUMBER:

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RESULT 10
US-10-284-060-26
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INFORMATION FOR SEQ ID NO:
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APPLICANT:
APPLICANT:
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APPLICATION NUMBER: US 60/007,255
FILING DATE: 06-NOV-1995
                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                  APPLICANT: Rocheleau, Thomas A
TITLE OF INVENTION: Insecticidal Protein Toxins from
TITLE OF INVENTION: Photorhabdus
                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: protein
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: 1
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APPLICATION NUMBER: US 0/
FILING DATE: 18-MAY-1993
ZIP: 46268
COMPUTER READABLE FORM:
                                                                                                                       NUMBER OF SEQUENCES:
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                                                        CITY: Indianapolis
                                                                                         ADDRESSEE:
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Similarity 70.0%;
7; Conservative
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Petell, James
Patig, Raymond
Schoonover, Sue
firench-Constant, Richard
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Roberts, Jean L
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Pred. No. 1.2e+02;
2; Mismatches 1
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APPLICANT: Penn, Sharron G.,
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: ANALYSIS OF GENE EXPRESSION IN BT 474
FILE REFERENCE: PB 0004 WO 8
CURRENT APPLICATION NUMBER: US/10/203,138A
CURRENT APPLICATION NUMBER: US/10/203,138A
CURRENT APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 26 May 2000 (26.05.00)
PRIOR FILING DATE: 26 May 2000 (26.05.00)
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 03 AUGUST 2000 (28.05.00)
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 03 AUGUST 2000 (03.08.00)
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 03 AUGUST 2000 (03.08.00)
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US-10-203-138A-10373
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Best Local Similarity
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FILING DATE: 28-FEB-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 28-AUG-1996
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/
PILING DATE: 06-NOV-1996
APPLICATION INMBER: US 08/
FILING DATE: 06-NOV-1996
ATTORNEY/AGENT INFORMATION:
NAME: BOTUCKI, ANDREA: 33651
REFERENCE/DOCKET NUMBER: 5-
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Molecular Dynamics, APPLICANT: Penn, Sharron G.
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APPLICATION NUMBER: US 08/395,497
FILING DATE: 28-FEB-1995
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
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LENGTH: 1189 amino aci
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US 60/236,359
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70.0%;
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Pred. No. 1.2e+02;
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RESULT 13
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US-09-724-676-49554
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                                                                                         Sequence 49554, Application US/09724676A
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT FILING NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 49554, Application US/09724676
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: PatentIn version 3.2
SEQ ID NO 49554
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 30 June 2000 (30.06.00)
NUMBER OF SEQ ID NOS: 15438
SOPTWARE: Molecular Dynamics Sequence Listing Engine SEQ ID NO 10373
LENGTH: 69
                                                          SEQ ID NO 49554
LENGTH: 99
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Best Local
TYPE; PRT
ORGANISM: Homo sapiens
-09-724-676A-49554
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 27 September 2000 (27.09.00)
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 21 September 2000 (21.09.00)
PRIOR APPLICATION NUMBER: US 09/608,408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 99
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
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OTHER INFORMATION:
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nes 7; Conserv
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87.5%;
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Pred. No.
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US-09-950-084-5149
; Sequence 5149, Application US/09950084
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CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR FILING DATE: 2001-05-19
PRIOR FILING DATE: 2000-05-19
PRIOR FILING DATE: 2000-05-19
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
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                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 09/417,811
PRIOR FILLING DATE: 1999-10-14
PRIOR APPLICATION NUMBER: US 09/353,718
PRIOR FILLING DATE: 1999-07-14
PRIOR APPLICATION NUMBER: US 09/266,557
PRIOR APPLICATION NUMBER: US 09/266,556
PRIOR PILLING DATE: 1999-03-11
PRIOR PILLING DATE: 1999-03-11
PRIOR APPLICATION NUMBER: US 09/266,556
PRIOR APPLICATION NUMBER: US 09/266,555
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: George H. Miller
APPLICANT: Roberta S. Hare
APPLICANT: Karen J. Shaw
TITLE OF INVENTION: Staphylococcus aureus Related Compositions and Methods
FILE REFERENCE: 1034/1263US2
CURRENT APPLICATION NUMBER: US/09/950,084
CURRENT FILING DATE: 2001-09-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: George H. Shimer, Jr.
APPLICANT: George H. Miller
APPLICANT: Roberta S. Hare
                                     PRIOR APPLICATION NUMBER: US 09/036,720
PRIOR FILING DATE: 1998-03-06
PRIOR APPLICATION NUMBER: US 09/036,338
PRIOR FILING DATE: 1998-03-06
                                                                                                                            PRIOR APPLICATION NUMBER: US 09/266,541
PRIOR FILING DATE: 1999-03-11
PRIOR APPLICATION NUMBER: US 09/037,934
PRIOR FILING DATE: 1998-03-10
                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 09/266,542 PRIOR FILING DATE: 1999-03-11
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Prior Application data removed - SEQ ID NOS: 7451
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77.8%;
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Pred. No. 12;
3; Mismatches
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; SEQ ID NO 5149
; LENGTH: 119
; TYPE: PRT
; ORGANISM: Staphylococcus aureus
US-09-950-084-5149
Query Match
Best Local Similarity 70.0%; Pred. No. 14;
Matches 7; Conservative 1; Mismatches 2; Indels 0; Gaps 0;
QY 1 KLKSQLVKRK 10
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Db 107 KLKSEKLEKRK 116
Search completed: February 8, 2003, 10:51:43
Job time: 16 secs
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Result
No.
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Maximum DB
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
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Match Length
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Gapop 10.0 , Gapext 0.5
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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                                               US-08-477-831C-38
US-08-90-178-3
US-08-90-178-3
US-08-91-654-3
US-08-91-344-4
US-08-477-831C-2
US-08-477-831C-11
US-08-801-344-4
US-09-458-599-4
US-09-18-5331-32
PCT-US91-05047-32
US-08-118-270-11
PCT-US93-08528-11
US-08-118-270-11
PCT-US93-08528-11
US-09-134-001C-3095
US-09-134-001C-4988
US-09-134-001C-4988
US-08-18-28-18-6
US-09-134-001C-4988
US-08-195-654-2
US-08-995-654-2
US-08-995-654-2
US-08-913-553-7
US-08-313-553-7
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Sequence 38, Appli
Sequence 3, Appli
Sequence 3, Appli
Sequence 2, Appli
Sequence 11, Appli
Sequence 4, Appli
Sequence 4, Appli
Sequence 592, Appli
Sequence 592, Appli
Sequence 32, Appli
Sequence 32, Appli
Sequence 31, Appli
Sequence 11, Appli
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ALIGNMENTS	PCT-US95-11856-7	US-08-309-419-7	US-08-309-420-7	US-08-462-040-48	US-08-460-971A-48	US-08-167-641C-48	US-08-460-890A-48	US-08-462-040-49	US-08-460-971A-49	US-08-167-641C-49	US-08-460-890A-49	US-09-122-171D-8	US-08-062-472B-26	US-08-477-831C-39	US-08-427-072-13	US-08-459-065-3	US-08-459-146-3	US-08-418-893D-24
	Sequence 7, Appli	Sequence 7, Appli	Sequence 7, Appli	Sequence 48, Appl	•	•	•	•	Sequence 49, Appl	•	Sequence 49, Appl	Sequence 8, Appli	Sequence 26, Appl	Sequence 39, Appl	Sequence 13, Appl	Sequence 3, Appli	Sequence 3, Appli	Sequence 24, Appl

RESULT 1 US-08-477-831C-38 US-08-477-831C-38 Sequence 38, Application US/08477831C Patent No. 6429291 CITY: NEW YORK STATE: NEW YORK COUNTRY: U.S.A. ZIP: 10020-1104 COMPUTER READABLE FORM: MEDIUM TYPE: Ploppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Rel. #1.0, ASCII CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/477,831C FILING DATE: 07-JUN-1995 CLASSIFICATION: 514 ATTORNEY/AGENT INFORMATION: NAME: PIERRI, MARGARET A. REGISTRATION NUMBER: 30,709 REFERENCE/DOCKET NUMBER: SIM-10 TELEFAX: 212-596-9090 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 32 amino acids GENERAL INFORMATION: APPLICANT: TURLEY APPLICANT: TURLEY, EVA A. APPLICANT: SHUWEN, ZHANG APPLICANT: ENTWISTLE, JOYCELYN TITLE OF INVENTION: HYALURONAN RECEPTOR PROTEIN NUMBER OF SEQUENCES: 60 CORRESPONDENCE ADDRESS: ADDRESSEE: FISH & NEAVE MOLECULE TYPE: DESCRIPTION: /desc = protein fragment TELECOMMUNICATION INFORMATION: TELEPHONE: 212-596-9000 TELEFAX: 212-596-9090 STRANDEDNESS: TOPOLOGY: lir TYPE: amino acids 1251 AVENUE linear single 93.5%; 얶 THE AMERICAS Score 43; DB Pred. No. 0.17 1; Mismatches Length 32

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KLKSQLVKRK 10

Local Similarity les 9; Conserv

Conservative

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RESULT 3
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                                                                  Sequence 3, Application US/08700178

Patent No. 5783669

Patent No. 5783669 5700912

GENERAL INFORMATION:
APPLICANT: Hawkins, Phillip R.
APPLICANT: Wilde, Craig G.
APPLICANT: Wilde, Craig G.
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: HYALURONAN RECEPTOR EXPRESSED IN HUMAN
TITLE OF INVENTION: UMBILICAL VEIN ENDOTHELIAL CELLS
NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                    CORRESPONDENCE ADDRESS
(ADDRESSEE: INCYTE P)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IMMEDIATE SOURCE:
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TOPOLOGY: lir
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33954
REFERENCE/DOCKET NUMBER: PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: HYALURONAN RECEPTOR EXPRESSED IN HUMAN TITLE OF INVENTION: UMBILICAL VEIN ENDOTHELIAL CELLS NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                424 KLRSQLVKKK 433
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CITY: Palo Alto
STATE: CA
                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LIBRARY: mouse
CLONE: GI 53979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/0
FILING DATE: 10-MAR-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE:
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Palo Alto
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               3174 Porter Drive
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                                    INCYTE PHARMACEUTICALS, INC
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US-08-995-654-3
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
**RNGTH: 477 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 3, Application US/08995654
Patent No. 6025138
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                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Hawkins, Phillip R.
APPLICANT: Wilde, Craig
APPLICANT: Seilhamer, Jeffrey
TITLE OF INVENTION: HYALUROWAN RECEPTOR EXPRESSED IN HUMAN
TITLE OF INVENTION: UMBILICAL VEIN ENDOTHELIAL CELS
NUMBER OF SEQUENCES: 3
                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: ISM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/700,178
FILING DATE: August 20, 1996
                               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IMMEDIATE SOURCE:
LIBRARY: mouse
CLONE: GI 53979
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FILING DATE: March 10, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REGISTRATION NUMBER: 36,749
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: Incyte Pharmaceuticals, Inc
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TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
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                                                                        APPLICATION NUMBER: US/08/995,654 FILING DATE: December 22, 1997
                                                                                                                                                                                                                                                                                  STREET: 3174 Por
CITY: Palo Alto
                                                                                                                                                                                                                             ZIP: 94304
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TOPOLOGY: linear
FILING DATE:
                 APPLICATION NUMBER:
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August 20, 1996
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                   08/700,178
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Pred. No. 2.1;
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US-08-477-831C-2
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Best Local Similarity yu.u
Thes 9; Conservative
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Best Local Similarity
                 Query Match
                                                                                                                                                                           TELEPHONE: 212-596-9000
TELEFAX: 212-596-9090
INFORMATION FOR SEQ ID NO: 2:
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INFORMATION FOR SEQ ID NO:
                                                              TOPOLOGY: 1in MOLECULE TYPE: DESCRIPTION:
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REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0028-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: PIERRI, MARGARET A.
REGISTRATION NUMBER: 30,709
REFERENCE/DOCKET NUMBER: SIM-10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-596-9000
                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Rel. #1.0, ASCII CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,831C FILING DATE: 07-JUN-1995
CLASSIFICATION: 514
APPLICATION: 514
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LENGTH: 477 amino acid
                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                              LENGTH: 600 LENGTH: 600 CONTROL OF TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Dol "
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: NEW YORK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
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T: ENTWISTLE, JOYCELYN
INVENTION: HYALURONAN RECEPTOR PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NEW YORK
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                                                                                                                                           606 amino acids
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/desc = "RHAMM I protein"
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93.5%;
90.0%;
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Pred. No. 2.1;
Score 43; DB 4; Length 606; Pred. No. 2.6;
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                                                                                                                                    Sequence 4, Application US/08801344
Patent No. 6087140
GENERAL INFORMATION:
APPLICANT: Cameron, Douglas C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 11, Application US/08477831C Patent No. 6429291
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Matches 9; Conserv
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Rel. #1.0, ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,831C
FILING DATE: 07-JUN-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: PIERRI, MARGARET A.
REGISTRATION NUMBER: 30,709
REFERENCE/DOCKET NUMBER: SIM-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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                                               APPLICANT: Shaw, Anita J.
APPLICANT: Altaras, Nedim B.
TITLE OF INVENTION: MICROBIAL PRODUCTION OF 1,2-PROPANEDIOL
TITLE OF INVENTION: FROM SUGAR
NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
DESCRIPTION:
CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 631 amino acids
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CORRESPONDENCE ADDRESS:
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APPLICANT: ENTWISTLE, JOYCELYN
TITLE OF INVENTION: HYALURONAN RECEPTOR PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
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TELEPHONE: 212-596-9090
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SHUWEN, ZHANG
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/desc = "RHAMM I-2a"
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Pred. No.
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MOLECULE TYPE: protein
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
GRGANISM: Rat lens aldose reductase
US-08-801-344-4
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Best Local Similarity
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NAME: Sara, Charles S.
REFERENCE/DOCKET NUMBER: 0982
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-831-2100
TELEPACE: 608-831-2106
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 316 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/09498599 Patent No. 6303352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATME: Sara, Charles S.
(REFERENCE/DOCKET NUMBER: 0982
TELECOMMUNICATION INFORMATION:
                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/498,599
                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Cameron, Douglas C.
APPLICANT: Shaw, Anita J.
APPLICANT: Altaras, Nedim E.
TITLE OF INVENTION: MICROSIAL PRODUCTION OF
TITLE OF INVENTION: 1,2-PROPANEDIOL FROM SUC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
COMPUTER: PC COMPATEN: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                      COUNTRY: U.S.A. ZIP: 53717-1914
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70.0%;
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Pred. No. 38;
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APPLICANT: Wang, Aljun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
ITILE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
ITILE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
ITILE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
ITILE REFERENCE: 210121.427C16
CURRENT APPLICATION NUMBER: US/09/605,785
CURRENT FILING DATE: 2000-06-27
NUMBER OF SEQ ID NOS: 835
SOFTWARE: FRSCSEQ for Windows Version 3.0
SEQ ID NO 592
LENGTH: 188
TYPE: PRT
ORGANISM: Homo sapien
US-09-605-785-592
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                                                                Query Match
Best Local S
Matches 7
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 316 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Xu, Jiangchun APPLICANT: Dillon, Davis
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
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HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Rat lens aldose reductase
178 LKAQLVKQK 186
                                                              Local Similarity hes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        62 KLKEQVVKRQ 71
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STRANDEDNESS: sir
TOPOLOGY: linear
                   2 LKSQLVKRK 10
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Vedvick, Thomas
Carter, Darrick
Li, Samuel
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Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas
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Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Henderson, Robert A. Kalos, Michael D.
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                                                                 Conservative
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                                                                               Score 34; DB 4;
Pred. No. 35;
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Pred. No. 38;
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RESULT 10
US-07-725-331-32
; Sequence 32, Application US/07725331
; Patent No. 5294605
; GENERAL INFORMATION:
; APPLICANT: Houghten, Richard

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PCT-US91-05047-32
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                                                                                                                                                                                                                                                          Sequence 32, Application PC/TUS9105047
GENERAL INFORMATION:
APPLICANT: Houghten, Richard
APPLICANT: Blondelle, Sylvie
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Best Local S
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 amino acids
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: Chicago
CITY: IL
TISA
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ADDRESSEE: Dressler, Goldsmith, Sutker, Shore ADDRESSEE: & Milnamow
                                                                                                                                                                                                         TITLE OF INVENTION: Amphiphilic Peptide Compositions and TITLE OF INVENTION: Analogues Thereof NUMBER OF SEQUENCES: 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Rel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ADDRESSEE: Dressler, Goldsmith, Sutker, Shore, ADDRESSEE: & Milnamow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 19-JUL-1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Blondelle, Sylvie
TITLE OF INVENTION: Amphiphilic Peptide Compositions and
TITLE OF INVENTION: Analogues Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION
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                                                                                                                    STREET: 180 No
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nes 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/07/725,331
                                                                        ZIP: 60601
                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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                                                                                                                                         180 North Stetson
                                                                                        USA
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3126165460
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Pred. No. 8.9;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3, Application US/08585595
Patent No. 5795761
                                        PILING DATE: 16-JAN-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/584
FILING DATE: 11-JAN-1996
ATTORNEY/AGENT INFORMATION:
NAME: AUERBACH, JEFFREY I.
REGISTRATION NUMBER: 32680
REFERENCE/DOCKET NUMBER: 6137
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 383-7451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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              TELEFAX: (202) 383-6610 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: ANDERSON, STEPHEN
TITLE OF INVENTION: IMPROVED METHODS FOR PRODUCING VITAMIN
NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
ADDRESSEE: HOWREY & SIMON
                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC COMPATIBLE
COMPUTER: IEM PC COMPATIBLE
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 18 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 19-JUL-1990
ATTORNEY/AGENT INFORMATION:
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SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                      CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
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                                                                                                                                                                                                                                                                                                                                                      ZIP: 20004
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TELEFAX: 3126165460
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                                                                               6137-0014 CIP
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Pred. No. 8.9;
3; Mismatches
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; STRAIN: HOMO SAPIENS
US-08-585-595-3
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                                                                                                                                           INFORMATION FOR SEQ ID NO: 11
SEQUENCE CHARACTERISTICS:
LENGTH: 355 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                      Query Match
Best Local Similarity
Matches 6; Conserv
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Patent No. 5508384
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Best Local &
                                                                                                                            MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/118,270
FILING DATE: 09-SEP-1993
PRIOR APPLICATION NUMBER: US 07/943,236
APPLICATION NUMBER: US 07/943,236
                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Murphy, Randall B.
APPLICANT: Schuster, David I.
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 34,033
REFERENCE/DOCKET NUMBER: MU
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                                                                                                  FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
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TYPE: amino acid
TOPOLOGY: 1455
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236 KTRSQITKRK 245
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419 Seventh Street, N.W., Suite 300
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                                                       Score 32; DB 1;
Pred. No. 1.4e+02;
2; Mismatches 2
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RESULT 15
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; ORGANISM: Staphylococcus epidermidis
US-09-134-001C-3095
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                                                                     CURRENT APPLICATION NUMBER: US/09/134,001C
CURRENT FILING DATE: 198-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1997-11-08
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 395
                                                                                                                                                                                                                                   Sequence 3095, Application US/09134001C
Patent No. 6380370
GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al
APPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC-007
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Best Local Similarity
Matches 6; Conserv
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GENERAL INFORMATION:
APPLICANT: New York University
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 355 amino acid
                                         LENGTH: 66
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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STRANDEDNESS: sin
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REGISTRATION NUMBER: 34,033
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419 Seventh Street, N.W., Suite 300
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Pred. No. 1.4e+02;
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Query Match 67.4%; Score 31; DB 4; Length 66;
Best Local Similarity 66.7%; Pred. No. 45;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
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Search completed: February 8, 2003, 10:44:05

Job time: 12.3333 secs

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seq28 (AA) ID seq28 AA preliminary pattern
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Number of sequences searched:
Number of sequence hits:
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Number of sequences searched:
Number of sequence hits:
Number of separate matches:
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GENPEPT, Release 131.0, Released on 30Aug2002, Formatted on 31Aug2002
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348,344,575
1,135,942
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ck: 3482 len: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <(R,K,H) (A,R,N,C,Q,G,H,I,L,K,M,F,P,S,T,W,Y,V) \{7,7\} (R,K,H) > (K) (R,Q,K,F,T) \{7\} (H) KRTRFRKQH
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                                                                                                                                                                                                                                                                                                            ! GB:V01492 bradykinin [Bos taurus] (ver 1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ! GB:AF384615 ribosomal protein 16 [Howittia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ! GB:S95530 zinc finger protein [Homo sapiens
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 Total finds:
Total length:
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05:55.83
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                                                                                                                                               ! GB:V01492 bradykinin [Bos tau]
                                                                                                                                                                           1 GB:S95530 zinc finger protein
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CAAOO584 Length: 9 February 11, 2003 12:16 Type: P Check: 3637 ...

1 KLIYILITVR

1 KLIYILITVR

1 KLIYILITVR

1 CAAO1163 Length: 9 February 11, 2003 12:16 Type: P Check: 3637 ...

1 KLIYILITVR

1 KLIYILITVR

1 KLIYILITVR

1 KLIYILITVR

1 AM50387 GB:AF384615 ribosomal protein 16 [Howittia trilocularis] (ver 1)

AAM50387 GB:AF384615 ribosomal protein 16 [Howittia trilocularis] (ver 1)

AAM50387 Length: 9 February 11, 2003 12:16 Type: P Check: 3482 ...

1 KRTRFRKQH

1 IAA SEQUENCE 1.0

AAB22067 GB:S95530 zinc finger protein [Homo sapiens] (ver 1)

AAB22067 Length: 9 February 11, 2003 12:16 Type: P Check: 3653 ...

1 RNQVTIYTK

1 AAB22067 GB:V01492 bradykinin [Bos taurus] (ver 1)

CAA24737 Length: 9 February 11, 2003 12:16 Type: P Check: 3472 ...

1 RPPGFSPFR
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ACCESSION
VERSION
KEYWORDS
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ORIGIN
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58
TTATACACAC AATGATATGT AAAAATAGTA TTCGACATAA
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                                                                                                                           GGAATGGAAT
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YTSEEGSHFMERCKVKKSPYGGWIVE"
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                                                                                                                                                                                                                                                                                                                                                                complement (2652.
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                             AATAGTATAT TGAATTCTTT
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                                                                                           GATTAATTTA CATAATACCT
                                                                                                                         CATCITIGI TITCITAGGI AACATAAITG
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he FPV,
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including
                               ATCATGTAGA
                                                            TGTAAAATCA
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the genes D7,
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1001 1451 1401 1301 1201 1151 1101 1051 1651 1601 1551 1351 1251 701 451 401 351 301 CTTAGGCCGT AAAAGATATA CATCTCAGGG ACAGGAGAAT ACCCGACGAA CCAAGGTATC TTGTAGAAGT AGAATTGACG TACCTTATTA TTTGGTAATA AGTATTATTA AGATCAATAG TCATATATTC ACATTTTTCT TGGAAATGTT AACGGCGACG TGACGGTATT ATACCTATAA TCGGTATAAG CTGTATATTC ACATATAAAT ATGGGGGAGT GGGTATTGTA AAATATTAAA ACTATCGGAA GGAGGAAGAG TTAAAAATAA AGAATCAATA TACCATTCAC ATATAGCGGA TAAAGTATAT ATCAAAAAGA CTTAGCGATA GACGAATTCA CATATCGCGT TTCGAAACAA ATAAAGAATA CCAGAGAACA ACAAAATATA TTAGAAAAAA ATAAAGATTG CATAAGTGAA AAATAAACAA CTATAAAAAC TTTTTTATAT CTTTATATCT ACTITAACGA ACGCGGTATA TCAAAAAAAG TCATAGTTAT GGATTTAGAC CAACAGGTAA TTATGGAGGG ACACGAAAAT TATATGGTAG GTCGGATGGT TACATGCATA CTGATTATAA TAAAGAAGCA GATACGACAC CGTGGCAACA TTGTTTTATT AAAAGCACCT TTGATAATAT TGATAATATA CAAAAAATCA AACTGGTAGC TAGTGTGTTT ACTTACCATA GAAGATAAGA GAAAGCGATG GTATAATAAC AGAAGCTTTT TAATTAAAAA AATATATTCA TATAATCGTT AACTTATTAG TATTGCTATA AAGTAAAATA ACTTCAGAGG AAGGATCACA TGGTAAAATC GTATATGTGC ATAGGATCGC GGGAACAACA GACTTCGGGA AAGCCAAGTC AACATAAAAT CGGCCGCTAA AAATTGAATG ACGAAGTAAG TATAACCCCT AAAAATGAAT CTAGGAGTGT TTACTGTTTC GAAGGTTCAT ATATGTGTGT GGGTGACGGA GAAGTTACTC CCCTATGTAC TTCTCTGCTT TATATATCTA AACATAGAAA AGTTTACGAT TCCGTACGGT GGATGGATAG GACAAGTGGG AAAGATGATC ATTATAAAAA TAAATTGCTC ATGCTATTTC ATTATCTCCT TCGTACCGGA TACTTGCATT TCAAAGAATT TATAGACCAA GAATACTTAC TTCAACAACT GAAATTAAAG CAGAAATATT AAGAACGTCT ACTATAGAAA GTAGTTACCT TAAGTATCCC CCAAACCATG AGGGTGCTGG ATTTCTCAAT TACTTCGTAG TATCAATGCC TTGAAGAGTT AAATATCATT AGCCGTCGAT TTTATGTATC TATGTATTAC ACACTAACAT TAAGATCATG ACTATATATA TAAATGGGAI TTTAATTATA TAGAATAGAT TTTTATGGAA TTATCAGATA GTGGAGATAT AAATCCTACA GTTAACTTAC GGTGGAAGAT CGAAGATCGA TGTATGGATA CTACATATAT AAGGAATAAA AGAAGATAAA AGTAGTGATA TCAAAAACTT CTCTATTCAG TGCTTTTAGT TATAAAAACA

2651 2601 2501 2451 2401 2301 2351 2251 2201 2151 2101 2051 TCTATCTAAC TGTTAATATA TAAATTAATT T TAATGAAAAC AATAGATGAC AAGTTTGTTT TATGCCATAG ATACAATAGT GATAAAGATT TTGAAGTTAT AGCCCTTTAC GTAGAAACAG ATCTTACGAG ATAATCTTCC CAGGTGGTAA AATAAAAGAC TTGGAAAGTA TAACTAATTG TTCTTGTTTT CAGAAATAGC ATTTACAAAA GATAAACGAC GGAAAATAAG ATATGAATAT GGCAAACTAC ATGCTAAATA TCCACTATCC GTAATAGGTA ATTAAATATA TAATAAACGC TGTAAGAACA TCCGCTAGCA ATAGTTAACA CATTTATAGA CGCAAGAGAT ATTAACAAAG ATTATTTGTA TACTAATGTA TAGACAAATA TTGAATAGAT TCATCCCTAA TAGAGAAATT AAAGGAATAT CTATCTGTAA GAACTGTTTT GTATACGGTT CTATATACGA CAGATTGATA TCTAGTAAGA GAAATAAAAG AAGAATTAAA TATCGATTCC TCTTATCTCG TAAGTTATAA GCTATCGCTT CCTAATAACT ATAATACAAA CCATATAGAT ACTCTTTAAG AAATATTCAA AGTATATGAG TAACATCGAA CGTGATATAT

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TITLE
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                              ACGTATAGCT TATTAAACAT GGATAGCATT TAGTAATATT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 2681)
Binns, M.K., Boursnell,
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db_xref="taxon:10261"
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AATAGTATAT TGAATTCTTT
                                                                                                  CATCITITGT TITCTTAGGT AACATAATTG
                                                                GATTAATTTA CATAATACCT
                                                                                                                                    2003 15:09
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1701 1951 1601 1551 1451 1251 1201 1151 1001 2001 1851 1751 1651 1501 1351 1301 1801 1401 1051 901 851 803 751 651 601 501 451 401 351 301 251 201 701 CTTAGGCCGT CTGTATATTC TGATAATATA CAAAAAATCA AACTGGTAGC TACCTTATTA GTTTATCAAA CATAAGTGAA AAATAAACAA CTATAAAAAC TAAGATCATG GAGTAGATAT AACAACTTGG GTAATTGTAG AGAATTGACG AAGATATTAG AGTATTATTA ATAAACCATC ATCAGATAAT ATACCTATAA TCGGTATAAG AAGAACGTCT TTTATGTATC AAAAGATATA ACACTAGTAT TCATATATTC CATCTCAGGG AGATGCAAAG CAGAATCTGA ACAGGAGAAT ACCCGACGAA AAAAAAAACC AACGGCGACG CCAAGGTATC TAAATCTTAC TTGTAGAAGT TTTGGTAATA AGATCAATAG TGTTATTACT TTCGATATAT ACATTTTTCT TGGAAATGTT TGTTACATAC TGACGGTATT ACATATAAAT ATGGGGGAGT CAAGTAATGA GGGTATTGTA TAGTGTGTTT TATAGACCAA CTCTATTCAG AAATATTAAA ACTTACCATA ACTATCGGAA GGAGGAAGAG TTAAAAATAA TACCATTCAC ATATAGCGGA TTCAACAACT TAAAGTATAT GTATAATAAC ATCAAAAAGA AGAAGCTTTT CTTAGCGATA GACGAATTCA CATATCGCGT TATGTATTAC TTCGAAACAA CCAGAGAACA TTTTTTATAT AGTTTACGAT ATGCTATTTC ACTATATATA CTTTATATCT TAATTAAAAA AATATATTCA ACTITAACGA TATAATCGTT AACTTATTAG ACGCGGTATA TATTGCTATA AAGTAAAATA TCAAAAAAAG TCATAGTTAT ACTTCAGAGG GGATTTAGAC TGGTAAAATC CAACAGGTAA GTATATGTGC TTATGGAGGG ACACGAAAAT TATATGGTAG AAGCCAAGTC AACATAAAAT GGTGGAAGAT GTCGGATGGT TACATGCATA CTGATTATAA AAAAATGAAT TAAAGAAGCA GAAGGTTCAT GATACGACAC GACAAGTGGG CGTGGCAACA TIGITITATI AMAGATGATC AAAAGCACCT TTGATAATAT TTATACACAC AATGATATGT AAAAATAGTA ACCTTTATAT AAAATACGAA AACATAGAAA GAAAGCGATG ATAAAGAATA ACAAAATATA TCCGTACGGT GGATGGATAG ATAGGATCGC GGGAACAACA GACTTCGGGA CGGCCGCTAA AAATTGAATG ACGAAGTAAG GGGTGACGGA GAAGTTACTC CCCTATGTAC TICICIGCII GAAGATAAGA GAAATTAAAG CTAGGAGTGT AGGGTGCTGG AGAAGATAAA TATATATCTA ATTATAAAAA TAAATTGCTC TACTTGCATT TCAAAGAATT GAATACTTAC AGAATCAATA CAGAAATATT ACTATAGAAA TTAGAAAAAA AAGGATCACA ATTATCTCCT GTAGTTACCT TAAGTATCCC AAATCCTACA TATAACCCCT ATATGTGTGT CCAAACCATG TCGTACCGGA TACTTCGTAG TATAAAAACA TATCAATGCC TTGAAGAGTT AAATATCATI ACACTAACAT ATAAAGATTG TAAATGGGAT TTTAATTATA ATTTCTCAAT TAGAATAGAT TTTTATGGAA TTATCAGATA GTGGAGATAI GTTAACTTAC TGCTTTTAGI TTACTGTTTC CTACATATAT AAGGAATAAA AGTAGTGATA TCAAAAACTT TTCGACATAA TTATTTATAA CGAAGATCGA TGTATGGATA AGGATCATTA AGCCGTCGAT

2651 2601 2551 2451 2351 2401 2301 2251 2201 2101 2151 2051 GATAAAGATT TIGAAGTTAT AGCCCTTTAC GTAGAAACAG ATCTTACGAG TCTATCTAAC TGTTAATATA TAAATTAATT T CATTTATAGA CGCAAGAGAT ATTAACAAAG ATTATTTGTA TACTAATGTA TAGACAAATA TTGAATAGAT TCATCCCTAA TAGAGAAATT AAAGGAATAT ATAATCTTCC CAGGTGGTAA AATAAAAGAC TTGGAAAGTA TAACTAATTG TAAGTTATAA GCTATCGCTT CCTAATAACT ATAATACAAA CCATATAGAT ACTCTTTAAG AAATATTCAA AGTATATGAG TAACATCGAA CGTGATATAT TAATGAAAAC AATAGATGAC AAGTTTGTTT TATGCCATAG ATACAATAGT ATATGAATAT GGCAAACTAC ATGCTAAATA TCCACTATCC GTAATAGGTA ATTAAATATA TAATAAACGC TGTAAGAACA TCCGCTAGCA ATAGTTAACA CTATCTGTAA GAACTGTTTT GTATACGGTT CTATATACGA CAGATTGATA TCTAGTAAGA GAAATAAAAG AAGAATTAAA TATCGATTCC TCTTATCTCG TTCTTGTTTT CAGAAATAGC ATTTACAAAA GATAAACGAC GGAAAATAAG

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AUTHORS
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AF384615

DEFINITION Howittia trilocularis ribosomal protein 16 (rpl16) gene, intron and partial cds; chloroplast gene for chloroplast product.

ACCESSION AF384615

VERSION AF384615.1 GI:21388904
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ATTCTGAGAG GTTATGAGTT AACCCGACAA CTGAACTAGA TATTGAAAGA
                                                                                                                                                                                                                                                                                                               GTCTATGAAG CATCTCATAA AGAGCAATGT AATAGAGCAT CAATAGAGAT
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                                          GGGTAGGATG GCGGAACAAA CCGGAGACCA CTTCATTTCT TTTTATTCTG
                                                                                        GAACCCGTGA ATGCAGAGAT
                                                                                                                                CATTGTAGAA TTCAGACCTA
                                                                                                                                                                            AMATTTTATT TAATTTTTT TATTAMATTA AMATATTAMA ACGAAGGCTC
                                                                                                                                                                                                                       CAAGAGCTTT AACTTAAGTC AATAAAGACT GAGAAGGTTG
                                                                                                                                                                                                                                                                     TCATCAAGAA ATCAATAATT
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Pfeil, B.E., Brubaker, C.L., Craven, L.A. and Crisp, M.D.

Phylogeny of Hibiscus and the tribe Hibisceae (Malvaceae) using chloroplast DNA sequences of ndhr and the rpl16 intron Syst. Bot. 27 (2), 333-350 (2002)

2 (bases 1 to 1138)

Pfeil, B.E., Brubaker, C.L., Craven, L.A. and Crisp, M.D.

Direct Submission

Submitted (23-MAY-2001) Center for Plant Biodiversity Research, CSIRO Plant Industry, GPO Box 1600, Canberra, ACT 2601, Australia Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chloroplast Howittia trilocularis
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Malvales; Malvaceae; Howittia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="sequence begins in the domain I 5' bulge"
<1111. .>1138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Howittia trilocularis"
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156 c 178 g 336 t
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                                                                                                                                AGCATTAATC GAGAAGCGAT
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751	GTTAAATTTT	TGTCGATCTG	GTTAAATTTT TGTCGATCTG AATCTGATAT GAATATGATA	GAATATGATA	AAAAACAAAA
801	TAAAGATTCG	TTATAACATT	ATAACAAAAC	TAAAGATTCG TTATAACATT ATAACAAAAC CAAGATAAGA CATTATCTAG	CATTATCTAG
851	AAATAGAATC	CGTGTTTAAT	TACTTATATA	AAATAGAATC CGTGTTTAAT TACTTATATA TATATCCAAT ATATATCCAA	ATATATCCAA
901	ACAAGATATA	CAAATTTCTA	ATAGATCAGA	ACAAGATATA CAAATTTCTA ATAGATCAGA TAAAATATCA AAGCAAAGAA	AAGCAAAGAA
951	TCCCAAGATT	CAATCTATTA	TTCATTAACT	TCCCAAGATT CAATCTATTA TTCATTAACT ACCTGTATAT CTTAAGAATA	CTTAAGAATA
1001	AAATAGTTTT	TTAGTCATTT	TTTTTTCGCG	AAATAGTTTT TTAGTCATTT TTTTTTCGCG AGGAGCTGGA	TGAGAAGAAA
1051	CTCTCATGTC	CAGTTCTGTA	GTAGAGATGG	CTCTCATGTC CAGTTCTGTA GTAGAGATGG AATTCATAAA CAACCATCAA	CAACCATCAA
1101	CTATAACCCA	AAAAGAACCA	CTATAACCCA AAAAGAACCA GATTTCGTAA ACAACATA	ACAACATA	

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ACCESSION
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                                       $95530 Length: 30 February 11, 2003 15:09 Type: N
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CGCAATCAGG TGACAATTTA TACCAAATGA
                                                                                                                                                                                                                                                                                                                                                                                                                                     GenBank staff at the National Library of Medicine created this entry (NCBI gibbsq 95530) from the original journal article. This sequence comes from Fig. 5a.

Map location: 11p13.

Map location in the mucleotides 9-10 has resulted in a complete deletion between nucleotides 9-10 has resulted in a complete deletion of zinc finger domain. The deleted nucleotides are corresponding to 190-415 in normal DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Maitland,N.J.
Inactivation of the remaining allele of the WT1 gene in a Wilms' tumour from a WAGR patient Oncogene 7 (4), 763-768 (1992) 92228507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 30)

Brown, K.W., Watson, J.E., Poirier, V., Mott, M.G., Berry, P.J. and
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Homo sapiens
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                                                                                                   12
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                                                                                                                                                                                    /gene="WT1"
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                                                                                                                                                                                                                                                                                                  note="zinc finger protein"
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CGCCAGAATG GATAATCCTG
                                     AGTGAAAACA
                                                                          ATGACCAGGA
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BOYLINE messenger RNA coding for kininogen (clo gives rise to bradykinin a vasoactine peptide. V01492 K01758

V01492 J GI:493
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nature 305 (5934), 545-549 (1983)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kitamura,N., Takagaki,Y., Furuto,S., Tanaka,T., Nawa,H.
Nakanishi,S.
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Mammalla; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
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                                                                                                                                                                                                                                                                                                                                                     /product="bradykinin"
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/db_xref="SWISS-PROT:P01045"
/translation="RPPGFSPFR"
a 672 c 667 g 845 t
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                                                                                                                                              GTTAAATCAT GAAATTAATC ACCATCCTTT
                                                                                                                                                                                                                        AGAGCAGAGC
                                                                        TGTATTTAAA
                                                                                                             CCAAGTTTAA
                                                                                                                                                                                     TIGGCICCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SENKSGNQFVLYRITEVARMDNPDTFYSLKYQIKEGDCPFQSNKTWQDCDYKDSAQAA
TQCTATVAKKGNMKESVAIQTCLITPAEGFYVTAQYECLGCVHPISTKSPDLEFVLR
YAIQYFNNNTSHSHLFDLKEVKRAQKQVVSGMYEVYNSIAQTNCSKESFSFITFDCK
SLSGOTGECTDKAHVDVKLAISSFSQKCDLYFGEDFLFFMVCVGCFKFIFDDSPDLE
EALNHSIAKLNAEHDGTFYFKIDTVKKATVQVVGGLKYSIVFIARETTCSKGSNEELT
KSCEINIHGQILHCDANVYVVFWEEKVYFTVNCQFLGQTSLMKRFPGFSFFRSVQVMK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                QTEPNSLAFKLISDFPETTSPKCPSRPWKPVNGVNPTVEMKESHDFDLVDALL"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HGHHRSHGLGHGHQKQHGLGHGHKHGHGHGKHKNKGKNNGKHYDWRTPYLASSYEDST
TSSAQTQEKTEETTLSSLAQPGVAITFPDFQDSDLIATVMPNTLPPHTESDDDWIPDI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Bos taurus"
/db_xref="taxon:9913"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           translation"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TEGSTTVSLPHSAMSPVQDEERDSGKEQGPTHGHGWDHGKQIKLHGLGLGHKHKHDQG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /protein_id="CAA24736.1"
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                                                                                                                                                                                                                                                                                                February 11,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note="Protein sequence is"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    db_xref="SWISS-PROT:P01045"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'product="kininogen"
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ACACATTTA TICCITGAAG TACCAAATCA
                                                                                                                                                                                   ACTGGCTCCC AATTCCTGGT
                                                                        GCTGTGGACG
                                                                                                           CCCAAGAGTC
                                                                                                                                                                                                                        CCAGAGGAGA CAGAGGCTGA
                                                                                                                                                                                                                                                             ACATCAGGAA TCGCCAGGGC
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                                                                                                           CTCTCAAGAA
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                                   TAACCGAGGT
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                                                                        AAAATACAAC
                                                                                                           ATCGACTGCA
                                                                                                                                              TCCTTTGTTC
                                                                                                                                                                                   GAAACTCCTA
                                                                                                                                                                                                                        AGCTAGTTAA
                                                                                                                                                                                                                                                             CCAAAAGCAC
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1701 1351 1301 1251 1151 1801 1751 1551 1501 1451 1201 1051 1001 1951 1851 1651 1601 1401 951 901 851 801 601 451 401 751 701 651 501 AAGACTCCCC CTCTGAACCA TACAACAAAA AAGCCAGTTA TTTCAGATTT GCAACTGTGA GCCACGGAAA TGGACATCAA CATAAGCATG GGGACCATGG GGAAGCACAA GGCCTCCGGG TTACCCTACT ATTCTACACT GAAGTAATGA ATTGAAGTAT TATTTCAAGA CAGAAATGTG GTACAGATAA GAAGTTAATT TGAAAGAAGT TGCCATCCAA CTCCAGCCGA CAAGAGAGGG TACAAGGACT AGGAGGGCGA TTTTGATCTT CIGGATCCCI AGCCAGGTGT TGCACAGACG TGGAGGACAC TGAAGAGCGG TGTCGGCTGC ATTCTTAACT ATGGGAACAC CGTCAATCTC TATGGATGAC ATAAAACTGT ACCATGAACA TTCAGAACAG GTTGATGCTC ATGGAGTGAA TCCAGAAACA GACATCCAGA TGCCTAATAC ACTACCACCT CACACAGAGA GTGATGATGA AGCCATTACC CAAGAGAAGA CCTATTTGGC ACATAAAAAC AAAGGAAAAA ACAATGGAAA AAGCAACATG GTCTTGGCCA AAAGCAAATA AAATTACATG GATTCAGGAA AAGAACAAGG CTGTAAGTCT TTTTTCACCT TTCCGATCAG GTCAACTGTC AACCACTTGG GTGATGCTAA AGAGCTGACC TCTATTGTGT TTGACACTGT GAAAAAAGCA ACAGTACAGG TTCCATCGCA AAGCTTAATG CCCAAACCTA TACCTGTTGA ACCTITATCC AGGGGAGGAT AGCACATGTA GATGTCAAGC CCAGACTGCA ACTCAATTGC ACAAACTAAT AAAAAGAGCC CAAAAACAGG TATTTTAACA ACAACACCAG TATCTACCAA GGGCCCCGTG GTGACAGCCC AATATGAAGT CTGCACAAGC CTGTCCTTTT CAAAGTAACA AAACTTGGCA GGACTGTGAC ACCAAGGTCA ACCTCCCCCA CAGAGCCAAA ACCCCACTCT TGTCTATGTG TCATAGCAAG TCTCCGTGGC TTCTTTAATT TCCAACTGTG TTTCCTGACT CAGAAGAGAC AAGTTCTTAT TGGGCACCAT AAGAGTTGTG AGTCCCTTTC TGCCACAGGA GAGCCCCGAC TGTCCCCATT CCTAGAGAGA AATAAATGGT AATGTCCTAG TAGCCTTGCA TAGGCAGCCA GAAATGAAAG TTCAGGACTC AACACTCTCT GAAGATAGCA TGGACATAAG AGAAGTCATG GCCTTGGCCT TGGCCATAAA ACCCACTCAT GGGCATGGCT GCCATGTCAC CTGTACAAGA TTCAAGTGAT GAAAACTGAA ACAGACCTCA CTCATGAAAA GIGCCTIGGG AGATCAATAT GGAAACCACA TGTTCTAAGG CAGAGCATGA CAGCCCAGAC TTTTTACCCC TAAGAATTTC AAGTGGTGAT TGTTCCAAGG TGGTGTCTGG TCATTCCCAC CTCTTTGATC TIGGAGCCIG AGTATGAGTG TATCCAGACC CAGTGCACAG GGGGTATTTC CTACATCCTC GCATTATGAT CATGGTCATG TGGAACCTTC CCTTGGCTGT TTTAAATTGA AGATCTCATT TCCCTAGCCC GTCTTGGCCA AGGAAAAAGT CTGGAGGAAG CCATGGTTTG TTCCTTCTCG AGGAATTTTC CGACCGTGGC AGTCTCATGA TCGCCCCTGC ACATGGTCAA TGGTAGGTGG ACTGGTGAAT ATGGAACTAT TTCTGAGATA TGCCTGATCA

2951 3151 3101 3051 3001 2901 2851 2801 2751 2701 2651 2601 2551 2501 2451 2401 2351 2251 2201 TGCAAATAAA GAACCAATGA TCTC GGAAAGAATA ATATGATTAT CTTTAATTTA CTGGTAATGT GAGGGAGGCA AATAATAAA ACCTCCAGAG GTCAAAGAGG AACAAGAGAG AAAGACAGAT AGCACAGAAA ATGGACAAAC TAATATACTT GTGGCCTACT TCAGTTGCTT ACACCCACAC TCACCTGTGG ATGCCCCATCA CTGTTTCTTC AAATTATTGG TGTCTTTCAC AGCCCCTGCT GTGCTTTTGC CTAGGGAAAT AGGGCTTTAT TATAGTTGCA TTTTCTTTTG CTATAACATT GAAGTATATG GACTTTTGCA TGTGATGCAA ATTTTGTTCA GAAATTTTAT GAGAAATTCT TTGTAGCCTT CCTGCTTGAG GAAATCTTTT TTTAATAATT GATAAGGGAA TATCATGAGA GCTCTGAGTT TAATAATAAG GATGGAAAGC AGAAGTAATA CAAAGGCTGG TTACTCATTA ACTGCGCTTT GCAATAGGGC TTTCAAATAG GAGTTTTCTG ATAATTCTCT TTCTGGGTGA GAGTGTTTCT TGTAAGTCAG TAATTTCTGT TGGCCAAAGG GAGGACAGAG AAAATAAATG AATTGACTTT CTGTTTCCAA ATAAGGGAAG TTAAGAGACT GAATGTCGAA CTTCTTAATG GAGGGAAAAA AAGATTCCTG ACATTTTGAA TAAACTGTAA CATCTGGTCC TTATGTGGCA TTCTGATAAT GAATGTGTAC CCTCAAATGT ACAATGTCAA TTTTCATACA GTGCAGAATT CTAACTCCTT TCTGAGGTTT CCTCCCAGGT TTTCTTAACT ATGCGTTTGA GCCACTGAGT TTTTATTTCA TTCAAAATAG TATCTGTACT TITGTTTGTA TGTTGCCAGC ATTAGACTGA GTAATCCTTA CTGTTCAATA AATGGGCTAG TAATCTCAAA CAGTTACTCC CATGTGACAA AGCTATTCTG

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! FINDPATTERNS on pir:* allowing 0 mismatches
Total finds:
Total length:
Total sequences:
CPU time:
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NBRF, Release
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96,134,422
283,224
52.87
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                                                                                                     73.0, Released on 16Aug2002, Formatted on 20Aug2002
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                                                                                                                                                                                                   < (R,K,H) (A,R,N,C,Q,G,H,I,L,K,M,F,P,S,T,W,Y,V) \{7,7\} (R,K,H) > \\ (R) (G,F,P,S) \{7\} (R) \\ RPPGFSPFR 
                                                                                                                                                                                                                                                                                                                                                          <(R,K,H) (A,R,N,C,Q,G,H,I,L,K,M,F,P,S,T,W,Y,V) \{7,7\} (R,K,H) > (R) (G,F,P,S) \{7\} (R) RPPGFSPFR
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(R)(G,L,F,P,T)\{7\}(R)
RPPGFTPLR
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RPPGFSPFR
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Total finds:
Total length:
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CPU time:
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! FINDPATTERNS on pir:* allowing 0 mismatches
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                                                            Databases searched:
NBRF, Release 73.0, Released on 16Aug2002, Formatted on 20Aug2002
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C;Accession: Abub''s R;Conlon, J.M.; Hicks, J.W.; Smith, D.D.
R;Conlon, J.M.; Hicks, J.W.; Smith, D.D.
Endocrinology 126, 985-991, 1990
A;Title: Isolation and biological activity of a novel kinin A;Title: Isolation and biological activity of a novel kinin ([Thr(6)]bradykinin) from the turtle, Pseudemys scripta.
([Thr(6)]bradykinin) A60579; MUID:90126625; PMID:2298179
                                                                                                                                                                                                                                !!AA_SEQUENCE 1.0
P1;A60579 - bradykinin-like peptide - slider turtle
C;Species: Pseudemys scripta (slider)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 18-Aug-2000
C;Accession: A60579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Anastasi, A.; Erspamer, V.; Bertaccini, G. Comp. Biochem. Physiol. A 14, 43-52, 1965
A;Title: Occurrence of bradykinin in the skin of Rana temporaria.
A;Reference number: A61363
A;Accession: A61363
A;Status: preliminary
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N;Alternate names: Thr-6-bradykinin
C;Species: Megascolia flavifrons (garden dagger wasp)
C;Date: 08-Mar-1989 #sequence_revision 08-Mar-1989 #text_change 18-Aug-2000
C;Accession: A26744
C;Accession: A26744
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A;Title: Threonine(6)-bradykinin in the venom of the wasp Colpa interrupta (F.) presynaptically blocks nicotinic synaptic transmission in the insect CNS. A;Reference number: A61057; MUID:91130217; PMID:1980872
A;Accession: A61057
                                                       A; Reference number: A60579; A; Accession: A60579
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Rana temporaria (common frog)
C;Date: 09-Sep-1994 #sequence_revision 09-Sep-1994 #text_change 18-Aug-2000
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Toxicon 25, 527-535, 1987
Toxicon 25, For the venom reservoirs of the A;Title: Two kinins isolated from an extract of the venom reservoirs of the solitary wasp Megascolia flavifrons.
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C;Species: Colpa interrupta
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 18-Aug-2000
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P1;A61363 - bradykinin -
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A; Accession: A26744
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;Molecule type: protein
;Residues: 1-9 <CON>
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C;Comment: This peptide increases aortic blood flow but, unlike bradykinin mammalian systems, does not lower blood pressure in the turtle. C;Superfamily: unassigned animal peptides C;Keywords: plasma !!AA_SEQUENCE 1.0 Length: 9 RPPGFTPFR ornitho-kinin -February 11, chicken 2003 12:15 Type: ש Check: 3478 in

C;Accession: B60246
R;Kimura, M.; Sueyoshi, T.; Morita, T.; Tanaka, K.; Iwanaga, Adv. Exp. Med. Biol. 247A, 359-367, 1989
A;Title: Ornitho-kininogen and ornitho-kinin: isolation, characteristics. C; Superfamily: unassigned A;Status: preliminary A;Molecule type: protein A;Residues: 1-9 <KIM> A;Reference number: A60246; MUID:90102072; PMID:2603803 C;Species: Gallus gallus (chicken)
C;Date: 11-Dec-1992 #sequence_revi chemical structure. B60246 Length: 9 #sequence_revision 11-Dec-1992 #text_change 18-Aug-2000 February 11, animal peptides 2003 12:15 Type: ש Check: 3526 characterization Ø and

RPPGFTPLR

ilAA SEQUENCE 1.0
F1;S65433 - bradykinin - horn fly (fragment)
C;Species: Haematobia irritans (horn fly)
C;Date: 28-Oct-1996 #sequence_revision 13-Ma
C;Accession: S65433 A;ACCEBBACH. A;ACCEBBACH. A;ACCEBBACH. A;Molecule type: protein
A;Residues: 1-9 <NAK>
A;Residues: bradykinin; hydroxyproline; skin
C;Keywords: bradykinin; hydroxyproline (Pro) #status experimental R;Nakajima, T.; Yasuhara, T.; Erspamer, G.F.; Visser, J. Experientia 35, 1133, 1979
A;Title: Occurrence of Hyp(3)-bradykinin in methanol extracts A;Reference number: S65431; MUID:96215437; PMID:8647080 A;Accession: S65433 A;Status: preliminary the South African leptodactylid frog Heleophryne purcelli. A;Reference number: A43065; MUID:80024576; PMID:488255 A;Accession: A43065 P1,A43065 - hydroxyproline-3-bradykinin - frog (Heleophryne purcelli)
C;Species: Heleophryne purcelli
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 07-May-1999
C;Accession: A43065 A; Molecule type: protein A; Residues: 1-9 < WIJ> A; Title: Cloning and characterisation of angiotensin-converting enzymodipteran species, Haematobia irritans exigua, and its expression in maturing male reproductive system. R; Wijffels, G.; Fitzgerald, C.; Gough, Willadsen, P. S65433 A43065 SEQUENCE 1.0 J. Biochem. 237, 414-423, 1996 the source is designated Length: 9 Length: 9 RPPGFSPFR February 11, February 11, as Haematobia irritans exigua 2003 2003 12:15 12:15 J.; Riding, 13-Mar-1997 #text_change 13-Mar-1997 Type: P Type: G.; Elvin, שי Check: 3472 Check: C.; Kemp, 3472 of the enzyme from the skin of the

RPPGFSPFR

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Total finds:
Total length:
Total sequences:
CPU time:
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SWISS-PROT, Release 40.3, Released on 9Aug2002, Formatted on 20Aug2002
SPTREMBL, Release 21.0, Released on 15Jun2002, Formatted on 28Jun2002
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784,472
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(R)(A,G,F,P,T)\{7\}(R)
RPAGFTPFR
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RPPGWSPFR
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Total finds: Total length: Total sequences: CPU time:	Databases searched: SWISS-PROT, SPTREMBL, Re	\\End of list	SP_OV:P83056	SP_OV:P83058	SP_OV:P83059	SP_OV:P83060	SP_OV:Q9PRJ4	1 < (!!SEQUENCE_LI
al finds: 5 1 length: 247,523,443 equences: 784,472 CPU time: 04:16.14	es searched: SWISS-PROT, Release 40.3, Released on 9Aug2002, Formatted on 20Aug2002 SPTREMBL, Release 21.0, Released on 15Jun2002, Formatted on 28Jun2002	list	056 ck: 3433 len: 9 finds: 1 ! P83056 bombina variegata (yell	058 ck: 3478 len: 9 finds: 1 ! P83058 bombina variegata (yell	059 ck: 3478 len: 9 finds: 1 ! P83059 bombina orientalis (ori	060 ck: 3472 len: 9 finds: 1 ! P83060 bombina orientalis	RJ4 ck: 3557 len: 9 finds: 1 ! Q9prj4 lepisosteus osseus (lon	$1 < (R,K,H) (A,R,N,C,Q,G,H,I,L,K,M,P,P,S,T,W,Y,V) \{7,7\} (R,K,H) >$	ISEQUENCE LIST 1.0 FINDPATTERNS on swp:* allowing 0 mismatches
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01-OCT-2001 (TrEMBLrel. 18, Created)
01-OCT-2001 (TrEMBLrel. 18, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
Bradykinin-like peptide ([Thr6]bradykinin).
Bombina orientalis (Oriental fire-bellied toad).
Bombina orientalis (Oriental fire-bellied toad).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Battachia; Anura; Archeobatrachia; Bombinatoridae; Bomb
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Q9PRJ4;
Q9PRJ4;
01-MAY-2000 (
            TISSUE-SKIN SECRETION;
Chen T.B., Orr D.F., Bjourson A.J., McClean S., Rac
"Cloning and post-translational processing of frog
Submitted (JUL-2001) to the SWISS-PROT data bank.
-!- SUBCELLULAR LOCATION: SECRETED.
                                                                                                                                                                                                                                                                                                                                  Vasodilator;
                                                                                                                                                                                                                                                                                                                                            Chen T.B., Orr D.F., Bjourson A.J., McClean S., Rao P.F. "Cloning and post-translational processing of frog skin submitted (JUL-2001) to the SWISS-PROT data bank.
-!- FUNCTION: VASODILATOR.
-!- SUBCELLULAR LOCATION: SECRETED.
-!- TISSUB SPECIFICITY: SKIN.
Vasodilator; Bradykinin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-OCT-2001 (TrEMBLrel.
01-OCT-2001 (TrEMBLrel.
01-OCT-2001 (TrEMBLrel.
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Conlon J.M., Platzack B., Marra L.E.,
"Isolation and biological activity of
of the phylogenetically ancient fish,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-MAY-2000 (TrEMBLrel.
01-MAY-2000 (TrEMBLrel.
01-JUN-2001 (TrEMBLrel.
                                                                                                    NCBI_TaxID=8346;
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Chen T.B., Orr D.F., B
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Amphibia; Batrachia; Anura; Archeobatrachia; Bombinatoridae;
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Bombina orientalis (Oriental fire-bellied toad).
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SEQUENCE 9 AA; 1099 MW;
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Eukaryota; Metazoa; Chordata;
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                                                                                EQUENCE,
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                                                                               SUBCELLULAR LOCATION,
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SPECIFICITY:
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17,
SKIN
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                                                                                TISSUE SPECIFICITY.
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roa skin kininogens.";
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SOCCEPTANT OCCUPANT O
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                                                             P83056
                                                                                                                                                 TISSUE-SKIN SECRETION;
Chen T.B., Orr D.F., Bjourson A.J., McClean S., Rao P.F.
"Cloning and post-translational processing of frog skin
Submitted (JUL-201) to the SWISS-PROT data bank.
-!- SUBCELLULAR LOCATION: SECRETED.
-!- TISSUE SPECIFICITY: SKIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-OCT-2001 (TrEMBLrel. 18, Created)
01-OCT-2001 (TrEMBLrel. 18, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
Bradykinin-like peptide ([Ala3,Thr6]bradykinin).
Bombina variegata (Yellow-bellied toad).
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01-OCT-2001 (TrEMBLrel. 18, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
Bradykinin-like peptide ([Thr6]bradykinin).
Bombina variegata (Yellow-bellied toad).
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Chen T.B., Orr D.F., Bjourson A.J., McClean S., Rao P.F., Shaw C.;
"Cloning and post-translational processing of frog skin kininogens.";
Submitted (JUL-2001) to the SWISS-PROT data bank.
                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleos
Amphibia; Batrachia; Anura; Archeobatrachia; Bombinatoridae;
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                                                                                                                                                                                                                                                                                                                                               SEQUENCE,
                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=8348;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -!- SUBCELLULAR LOCATION: SEC-!- TISSUE SPECIFICITY: SKIN SEQUENCE 9 AA; 1074 MW; :
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                                                                                                                       SPECIFICITY:
9 AA; 1048
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GenCore version 5.1.3 Coppright (c) 1993 - 2003 Compugen Ltd

OM protein - protein search, using sw model

February 8, 2003, 10:57:01; ; Search time 14 Seconds (without alignments) 75.534 Million cell updates/sec

Run

Sequence: Perfect score: Title: US-09-685-010-30 54 1 KQKIKHVVKLK 11

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0

283224 seqs, 96134422 residues

Searched:

Total number of hits satisfying chosen parameters:

1326

Minimum Maximum DB seg length: 0 seg length: 11

Post-processing: Minimum Match Maximum Match Listing first 80

45 summaries

Database

PIR_73:*
1: pir1:*
2: pir2:*
3: pir3:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16,	15	14	13	12	11	10	9	8	7	6	5	4	3	2	_	No.	Result
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	ribosomal protein	cytokeratin 4 - bo	phosphoenolpyruvat	polyphosphate-gluc	bursin - chicken	pyridoxal kinase (endo-1,4-beta-xyla		C 3.4.25.1 proteas	neomyosuppressin -		x pr	gene NF2 protein -	118K stomach cance	peptidyl-dipeptida	ribosomal protein	neuropeptide TE-6	cytochrome-c oxida	beta-galactosidase	probable secreted	\sim	hypothetical prote	cytochrome-c oxida	ite	virG protein - Agr	$\overline{}$	-	ane protein	Description	

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hydrogensulfite re	alcohol dehydrogen	ELAV-like neuronal	unidentified 6.0/1	glyceraldehyde-3-p	monodehydroascorba	small nuclear ribo	bothropstoxin - ja	FMRFamide-like pro	24K protein 4302 -	PV1 protein - huma	neurokinin A - Atl	neurokinin A - rai	hypothetical prote	thrombospondin 2 -	ubiquinol-cytochro

ALIGNMENTS

C;Accession: A33917
R;Simmer, J.P.; Kelly, R.E.; Scully, J.L.; Grayson, D.R.; Rinker Jr., A.G.; Bergh, Proc. Natl. Acad. Sci. U.S.A. 86, 4382-4386, 1989
A;Title: Mammalian aspartate transcarbamylase (ATCase): sequence of the ATCase doma A;Reference number: A33917; MUID:89282776; PMID:2543974
A;Accession: A33917 A; Molecule type: mRNA A; Residues: 1-7 <BRI>A; Cross-references: GB:Z14976; NID:g311650; PIDN:CAA78699.1; PID:g584083 C; Comment: This virus is coronavirus related to human coronavirus 229E. C; Keywords: membrane protein C:Accession: PQ0663

C:Accession: PQ0663

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C:Accession: PQ0663

C:Accession: PQ0663 A;Cross-references: GB:M23652 C;Superfamily: rudimentary enzyme; aspartate/ornithine carbamoyltransferase homology; arbamoyl-phosphate synthase (glutamine-hydrolyzing) large chain homology; carbamoyl-pc;Keywords: hydrolase dihydroorotase (EC 3.5.2.3) - Chinese hamster (fragment)
C;Species: Cricetulus griseus (Chinese hamster)
C;Date: 09-Mar-1990 #sequence_revision 09-Mar-1990 #text_change 07-Nov-1997 membrane protein - porcine epidemic diarrhea virus (isolate Belgian CV777) (fragment) C;Species: porcine epidemic diarrhea virus C;Species: 14-Unl-1994 #sequence_revision 14-Jul-1994 #text_change 08-Oct-1999 A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-11 <SIM> Вþ δÃ Query Match Best Local : Query Match Best Local Matches Matches :|: |:| 1 EKVLHLV 7 2 QKIKHVV 8 l Similarity 3; Conserva h 38.9%; Similarity 42.9%; 3; Conservative Conservative 38.9%; Score 21; DB 2; Le Pred. No. 9.2e+02; Pred. No. 9.2e+02; Pred. No. 9.2e+02; ω •• Score 21; DB 2; Pred. No. 2.8e+05; 3; 'Mismatches 1 Length 11; Length 7; Indels Indels sequence of the ATCase domain 0; 0 Gaps Gaps 0; 0

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hydrogensulfite reductase (EC 1.8.99.3) chain 1 - Desulfovibrio thermophilus (fragment) N;Alternate names: bisulfite reductase; desulfofuscidin C;Species: Desulfovibrio thermophilus C;Species: Desulfovibrio thermophilus C;Date: 19-Mar-1997 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998 C;Accession: S11024 R;Fauque, G.; Lino, A.R.; Czechowski, M.; Kang, L.; DerVartanian, D.V.; Moura, J.J.G.; Biochim. Biophys. Acta 1040, 112-118, 1990
                                                                                                                        A; Molecule type: protein A; Residues: 1-6 <FAU> C; Keywords: oxidoreductase
                                                                                                                                                                                          Biochim. Biophys. Acta 1040, 112-118, 1990
A;Tltle: Purification and characterization of bisulfite reductase (desulfofuscidin) from A;Reference number: S11024; MUID:90335276; PMID:2165817
A;Accession: S11024
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J. Mol. Biol. 215, 537-547, 1990
A;Title: Binding of the regulatory protein VirG to the phased
A;Reference number: S13224; MUID:91039316; PMID:2231718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 virG protein - Agrobacterium sp. (fragment)
C;Species: Agrobacterium sp.
C;Date: 19-Mar-1997 #sequence_revision 01-Aug-1997
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A;Accession: A28709
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-7 <OLS>
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R;Olsen, D.B.; Hepburn, T.W.; Moos, M.;
Biochemistry 27, 2229-2234, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Species: Bacillus cereus
C;Date: 22-Aug-1988 #sequence_revision
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                   A;Molecule type: protein A;Molecule type: protein A;Residues: 1-10 <LET> C:Comment: Complex I, mitochondrial NADH-ubiquiquinone reductase, ranging from 5K to 75K.
                                                                                         R; Leterme, S.; Boutry, M.
Plant Physiol. 102, 435-443, 1993
A;Title: Purification and preliminary characterization of mitochondrial complex I (NA A;Reference number: PQ0775; MUID:94151437; PMID:8108509
A;Accession: PQ0783
                                                                                                                                                                                                         NADH2 dehydrogenase (EC 1.6.99.3) 30K chain - fava bean mitochondrion N;Alternate names: complex I 30K chain; NADH-ubiquinone reductase 30K C;Species: mitochondrion Vicia faba (fava bean) C;Date: 03-May-1994 #sequence_revision 07-Oct-1994 #text_change 03-Jun C;Accession: PQ0783
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PQ0783
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of

03-Jun-2002

(fragment) chain

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C;Accession: S43631
R;Freund, R.; Kadenbach, B.
Eur. J. Blochem. 221, 1111-1116, 1994
A;Title: Identification of tissue-specific
A;Reference number: S43624; MUID:94237150;
                                                                                                                                                                                                                                                                                                    hypothetical protein - parsley (2.5 Species: Petroselinum crispum (parsley) (2.5 Species: Petroselinum crispum (parsley) (2.5 Sep-1999 #sequence_revision 20-Sep-1999 #text_change 20-Sep-1999 (2.6 Sep-1996) (2.6 Sep-1996)
                                                                                                                                                                                                                                                                                                                                                                          RESULT
                                                                                                                                                                                                               A; Title: Functional analysis of a light-responsive plant A; Reference number: Z18259; MUID:95128172; PMID:7827494 A; Accession: T14906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytochrome-c oxidase (EC 1.9.3.1) chain VIIa, cardiac - rainbow trout (frag
C;Species: Oncorhynchus mykiss (rainbow trout)
C;Date: 20-Oct-1994 #sequence_revision 01-Nov-1996 #text_change 16-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT
S43631
                                                                                                                                        A; Cross-references: EMBL:S75395;
                                                                                                                                                         A; Molecule type: DNA
A; Residues: 1-8 <FEL>
                                                                                                                                                                                                                                                                 R;Feldbrugge, M.; Sprenger, M.; Dinkelbach, M.; Yazaki, Plant Cell 6, 1607-1621, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Keywords:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Genome: nuclear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C; Genetics:
                                                                                                                                                                                            A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Note:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: protein A; Residues: 1-10 <FRE>
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                                                                      Matches
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Best Local
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                                   4 IKHVVKL 10
1 MKHTLTI 7
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                                                                    Similarity 2; Conserv
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                                                                      Conservative
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                                                                                     29.6%;
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                                                                    Score 16; DB
Pred. No. 2.8e
3; Mismatches
                                                                                                                                        NID:g913201; PID:e194245
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Pred. No.
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PMID:8181469
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                                                                лв <u>г</u>
1.8e+05;
2;
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                                                                                                     Length 8;
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Biochem. J. 312, 281-200, 1553
A;Title: Catalysis by the large subunit of
A:Reference number: S59625; MUID:96077156;
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S59625
                                                                                                        RESULT
S65385
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C;Date: 20-Jul-1996 #sequence_revision
C;Accession: S59625
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A; Reference number: S33518
A; Accession: S33519
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A; Genome: mitochondrion
C; Keywords: electron tr
A; Reference number: A; Accession: S65385
                                                                                                                                                                                                                                                      A; Residues:
                                                                                                                                                                                                                                                                  A; Molecule
                                                                                                                                                                                                                                                                           A; Status: preliminary
                                                                                                                                                                                                                                                                                         A; Reference number: A; Accession: S59625
                                                                                                                                                                                                                                                                                                                                                                                      beta-galactosidase alpha chain - Escherichia coli (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Cross-references: EMBL: Z22875;
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A; Residues: 1-11 <BOY>
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2; Conserv
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40.0%;
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Pred. No. 6.7e+03;
1; Mismatches 0
                                                                                                                                                                                                                 Score 15; DB 2; Pred. No. 1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 16; DB 2;
Pred. No. 7.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                              13-Mar-1997
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                                                                     20-Feb-1998 #text_change 07-May-1999
                                                                                                                                                                                                                                                                                                   the second beta-galactosidase of PMID:7492325
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                                             U.; von
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                     Enzymic properties and amino-term
                                                                                                                                                                                                      Indels
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C; Date: 100784
C; Accession: JH0784
C; Accession: JH0784
R; Smart, D.; Shaw, C.; Curry, W.J.; Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, L.; Hallow, Johnston, C.F.; Thim, L.; Hallow, Johnston, Johnston, C.F.; Thim, L.; Hallow, Johnston, Johnston, C.F.; Thim, L.; Hallow, Johnston, Joh
RESULT 13
S78764
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ribosomal protein MRP-S23, mitochondrial - bovine (fragment) C;Species: Bos primigenius taurus (cattle) C;Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change C;Accession: S78764
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C;Species: Ascaris suum (pig roundworm)
C;Date: 10-Jun-1993 #sequence_revision 10-Jun-1993 #text_change 28-Apr-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              submitted to the Protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: protein A; Residues: 1-6 < SMA>
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C;Keywords: oxidoreductase
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                                                                                                                                                                                                                                                       RESULT 14
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40.0%;
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75.0%;
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Pred. No. 2.86
3; Mismatches
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Pred. No.
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Pred. No. 2.
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2.8e+05;
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2.8e+05;
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1e+04;
                                                                                                                                             #text_change 07-May-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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                               I-converting
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                                  enzyme inhibitory
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118K stomach cancer antigen - human (fragment)
C:Species: Homo sapiens (man)
C:Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 17-Mar-1999
C:Accession: A60356
R:Shiraishi, Y.
Int. J. Cancer 45, 783-787, 1990
A:Title: Western blotting analysis for malignant lymphoma and stomach cancer antigens francescent comber: A60356; MUID:90216080; PMID:2323853
A:Accession: A60356
A:Molecule type: protein
A:Residues: 1-9 <SHI>C:Keywords: glycoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A:Accession: JN0859
A:Accession: JN0859
A:Molecule type: protein
A:Residues: 1-7 </ANT>
A:Residues: 1-7 </ANT>
A:Experimental source: Intestine
C:Comment: The carboxyl-terminus is essential for the protein's expression of angiotensic;Superfamily: bradykinin-potentiating peptide
C:Superfamily: bradykinin-converting enzyme inhibitor
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Search completed: February 8, 2003, 10:59:06 Job time: 15 secs
                                                                                                                                                                  Query Match 25.9%;
Best Local Similarity 42.9%;
Matches 3; Conservative
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Best Local S
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Local Similarity 60.0%;
hes 3; Conservative
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2 VAKLE 6
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Pred. No. 2.8e+05;
1; Mismatches 1; Indels
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Pred. No. 2.8e+05;
2; Mismatches 2;
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Minimum DB seq length: 0 Maximum DB seq length: 11
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Maximum Match
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                          Score
      protein search, using sw model
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54
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   10:
11:
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13:
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44.4
44.4
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Listing first 45 summaries
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/cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US07_NEW_MPUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
2: /cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
3: /cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US00_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US00_PUBCOMB.pep:*
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     2003,
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) US-09-984-056-92

) US-09-984-057-69

) US-09-834-765-258

) US-09-834-765-325

) US-09-834-765-643

) US-09-872-832-41

US-09-872-832-41

US-09-834-765-284

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US-10-012-034A-88

US-09-878-603-20

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(without alignments)
22.169 Million cell updates/sec
  Sequence 46, Sequence 20, Sequence 16,
                                                                                                                                                                                                                                                       Description
                                    Sequence 92, Appl
Sequence 69, Appl
Sequence 69, Appl
Sequence 258, App
Sequence 325, App
Sequence 38, Appl
Sequence 38, Appl
Sequence 284, Appl
Sequence 284, Appl
Sequence 284, Appl
Sequence 666, App
Sequence 666, App
Sequence 72, Appl
Sequence 72, Appl
Sequence 72, Appl
Sequence 72, Appl
Sequence 72, Appl
Sequence 73, Appl
Sequence 72, Appl
Sequence 72, Appl
Appl
Appl
Appl
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21	21	21	21	21	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22
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10	9	9	10	9	12	12	12	12	12	12	12	10	10	10	10	10	10	12	12	12	10	10	10	10	10
US-09-826-177-32	US-09-805-301-3	US-09-878-603-23	US-09-942-121-7	US-09-933-999A-20	US-10-012-034A-123	US-10-012-034A-112	US-10-012-034A-111	US-10-012-034A-109	US-10-012-034A-104	US-10-012-034A-102	US-10-012-034A-57	US-09-734-520-112	US-09-734-520-111	US-09-734-520-109	US-09-734-520-104	US-09-734-520-102	US-09-734-520-57	-10	US-10-012-034A-99	US-10-012-034A-98	US-09-984-057-3	US-09-984-056-3	US-09-734-520-113		US-09-734-520-98
Sequence 32, Appl	Sequence 3, Appli	Sequence 23, Appl	Sequence 7, Appli	Sequence 20, Appl	-	11	Sequence 111, App	Sequence 109, App	Sequence 104, App	102	57,		Sequence 111, App	109	Sequence 104, App	•		113,	99,	Sequence 98, Appl	Sequence 3, Appli	Sequence 3, Appli	Sequence 113, App		Sequence 98, Appl

ALIGNMENTS

RESULT 1
US-09-984-056-92
; Sequence 92, Application US/09984056
; Patent No. US20020120106A1

GENERAL INFORMATION:

APPLICANT: BOGOCH, SAMUEL APPLICANT: BOGOCH, ELENO

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RESULT 2
US-09-984-056-69
; Sequence 69, Application US/09984056
; Patent No. US20020120106A1
                                                                                                                                                                                                                                                                                                                           ; TYPE: PRT; ORGANISM: Bacillus anthracis US-09-984-056-92
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CURRENT FILING DATE: 2001-10-26
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR FILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: 09/146,755
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 1998-09-04
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: 08/198,139
PRIOR APPLICATION NUMBER: 08/198,139
PRIOR FILING DATE: 1994-02-17
NUMBER OF SEQ ID NOS: 103
SOFTWARE: PATENTIN 2:1
                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn
SEQ ID NO 92
LENGTH: 9
                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: BOGOCH, ELENORE S.
TITLE OF INVENTION: ANTHRAX AND SMALL POX REPLIKINS AND METHODS OF USE
FILE REFERENCE: 09425-46903
                                                                                                                                              1 KHIVKIE 7
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                                                                                                                                                                                                                                       4; Conservative
                                                                                                                                                                                                                                                            51.9%;
57.1%;
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Pred. No. 1.1e+05;
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CURRENT APPLICATION NUMBER: US/09/984,057

CURRENT FILING DATE: 2001-10-26

PRIOR APPLICATION NUMBER: 60/303,396

PRIOR FILING DATE: 2001-07-09

PRIOR APPLICATION NUMBER: 60/278,761

PRIOR FILING DATE: 2001-03-27

PRIOR APPLICATION NUMBER: 09/146,755

PRIOR APPLICATION NUMBER: 09/146,755

PRIOR APPLICATION NUMBER: 09/817,144

PRIOR APPLICATION NUMBER: 09/817,144
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US-09-984-056-69
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CURRENT FILING DATE: 2001-10-26
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR FILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: 09/146,755
PRIOR FILING DATE: 1998-09-04
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR FILING DATE: 1908-09-04
PRIOR FILING DATE: 2001-03-27
                                                                                  Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 90
SOFTWARE: PatentIn 2.1
SEQ ID NO 69
LENGTH: 8
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PRIOR FILING DATE: 1994-02-17
NUMBER OF SEQ ID NOS: 103
SOFTWARE: Patentin 2.1
SEQ ID NO 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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Patent No. US20020151677A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 44.4%;
Best Local Similarity 71.4%;
Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: BOGOCH, ELENORE S.
TITLE OF INVENTION: REPLIKINS AND METHODS OF IDENTIFYING
TITLE OF INVENTION: REPLIKIN-CONTAINING SEQUENCES
                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: 08/198,139
PRIOR FILING DATE: 1994-02-17
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TITLE OF INVENTION: ANTHRAX AND SMALL POX REPLIKINS AND METHODS OF USE FILE REFERENCE: 09425-46903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: BOGOCH, SAMUEL APPLICANT: BOGOCH, ELENC
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KKKHTVK 7
                                         KIKHVVK 9
                                                                                         Conservative
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                                                                                                          44.48;
71.48;
                                                                                       Score 24; DB 10;
Pred. No. 1.1e+05;
0; Mismatches 2
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Pred. No. 1.1e+05;
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                                                                                                                                Length 8;
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US-09-834-765-258
                                                                          Query Match
Best Local Similarity
Matches 4; Conserv
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 325
LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 325, Application US/09834765
Patent No. US20020055478A1
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Best Local :
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LENGTH: 9
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APPLICANT: Steve Chappell Mitchell
APPLICANT: Daniel E.H. Afar
APPLICANT: Daniel E.H. Afar
APPLICANT: Daniel E.H. Afar
APPLICANT: AND Jakobovits
TITLE OF INVENTION: GTP-BINDING PROTEIN USEFUL IN TREATMENT
TITLE OF INVENTION: AND DETECTION OF CANCER
FILE REFERENCE: 129.6USU1
CURRENT APPLICATION NUMBER: US/09/834,765
CURRENT FILING DATE: 2001-09-21
CURRENT FILING DATE: 2001-09-21
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APPLICANT: Daniel E.H. Afar
APPLICANT: Daniel E.H. Afar
APPLICANT: Daniel E.H. Afar
APPLICANT: Aya Jakobovits
TITLE OF INVENTION: GTP-BINDING PROTEIN USEFUL IN TREATMENT
TITLE OF INVENTION: AND DETECTION OF CANCER
FILE REFERENCE: 129.6USU1
CURRENT APPLICATION NUMBER: US/09/834,765
CURRENT FILING DATE: 2001-09-21
PRIOR APPLICATION NUMBER: 60/197,647
PRIOR APPLICATION NUMBER: 60/197,647
PRIOR FILING DATE: 2000-04-12
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APPLICANT: Pia M. Challita-Eid
APPLICANT: Arthur B. Raitano
APPLICANT: Steve Chappell Mitcl
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PRIOR FILING DATE: 2000-04-12
NUMBER OF SEQ ID NOS: 770
SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Pia M. Challita-Eid
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les 4; Conservative
2 QPYEHVIK 9
                                       2 QKIKHVVK 9
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                                                                                      Conservative
                                                                                                          44.4%;
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                                                                                   Score 24; DB 10;
Pred. No. 1.1e+05;
2; Mismatches 2
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Pred. No. 1.1e+05;
2; Mismatches 2
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US-09-834-765-643

Sequence 643, Application US/09834765

RESULT 6

Patelit No.

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RESULT 8
US-09-872-832-41
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Sequence 41, Application US/09872832
Patent No. US20020131960A1
GENERAL INFORMATION:
APPLICANT: Memorial Sloan-Kettering Cancer Center
TITLE OF INVENTION: ARPIFICIAL ANTIGEN PRESENTING
FILE REFERENCE: 830002-2003.1
CURRENT APPLICATION NUMBER: US/09/872,832
CURRENT FILING DATE: 2001-06-01
PRIOR APPLICATION NUMBER: 60/209,157
PRIOR APPLICATION NUMBER: 60/209,157
PRIOR FILING DATE: 2000-02-06
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Patent No. US20020131960A1
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Best Local :
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TITLE OF INVENTION: ARTIFICIAL ANTIGEN PRESENTING CELLS AND METHODS OF USE THEREOF
FILE REFERENCE: 830002-2003.1
CURRENT APPLICATION NUMBER: US/09/872,832
CURRENT FILING DATE: 2001-06-01
PRIOR APPLICATION NUMBER: 60/209,157
PRIOR FILING DATE: 2000-02-06
NUMBER OF SEQ ID NOS: 49
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CURRENT FILING DATE: 2001-09-21
PRIOR APPLICATION NUMBER: 60/197,647
PRIOR FILING DATE: 2000-04-12
NUMBER OF SEQ ID NOS: 770
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TYPE: PRT
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Local Similarity 50.0%;
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Arthur B. Raitano
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50.08;
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Pred. No. 1.1e+05;
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Pred. No. 1.1e+05;
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                                                                                                    CELLS AND METHODS OF USE THEREOF
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; ORGANISM: Homo sapiens US-09-834-765-89
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 4; Conserv
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SOFTWARE: Patentin version 3.0
SEQ ID NO 41
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NUMBER OF SEQ ID NOS: 770
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 284
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                                               APPLICANT: Aya Jakobovits
TITLE OF INVENTION: GTP-BINDING PROTEIN USEFUL IN TREATMENT
TITLE OF INVENTION: AND DETECTION OF CANCER
FILE REFERENCE: 129.6USU1
CURRENT APPLICATION NUMBER: US/09/834,765
CURRENT FILING DATE: 2001-09-21
PRIOR APPLICATION NUMBER: 60/197,647
PRIOR FILING DATE: 2000-04-12
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TYPE: PRT
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APPLICANT: Pia M. Challita-Eid
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TYPE: PRT
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Arthur B. Raitano
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Pred. No.
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TITLE OF INVENTION: GTP-BINDING PROTEIN USEFUL IN TREATMENT
TITLE OF INVENTION: AND DETECTION OF CANCER
FILE REFERENCE: 129 GUSU1
CURRENT APPLICATION NUMBER: US/09/834,765
CURRENT FILING DATE: 2001-09-21
PRIOR APPLICATION NUMBER: 60/197,647
PRIOR APPLICATION NUMBER: 60/197,647
PRIOR FILING DATE: 2000-04-12
NUMBER OF SEQ ID NOS: 770
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 386
LENGTH: 10
ORGANISM: Homo Sapiens
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Watches 4; Conserve
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-834-765-284
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APPLICANT: Mary Faris
APPLICANT: Pia M. Challita-Eid
APPLICANT: Arthur B. Raitano
APPLICANT: Steve Chappell Mitchell
APPLICANT: Daniel E.H. Afar
                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ 15 NO 666
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                                                                                                          APPLICANT: AYA JAKOBOVÍTS
TITLE OF INVENTION: GTP-BINDING PROTEIN USEFUL IN TREATMENT
TITLE OF INVENTION: AND DETECTION OF CANCER
FILE REFERENCE: 129.6USU1
CURRENT APPLICATION NUMBER: US/09/834,765
CURRENT FILING DATE: 2001-09-21
PRIOR APPLICATION NUMBER: 60/197,647
PRIOR APPLICATION NUMBER: 60/197,647
PRIOR FILING DATE: 2000-04-12
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APPLICANT: Pla M. Challita-Eid
APPLICANT: Arthur B. Raitano
APPLICANT: Steve Chappell Mitchell
APPLICANT: Daniel E.H. Afar
                                                                                              NUMBER OF SEQ ID NOS: 770
                     TYPE: PRT
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ORGANISM: Homo sapiens
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20020055478A1
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Pred. No.
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; NAME/KEY: AMIDATION
; LOCATION: (0)...(11)
US-09-734-520-88
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US-09-734-520-88
; Sequence 88, Application US/09734520
; Patent No. US20020115173A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/734,520
CURRENT FILING DATE: 2000-12-11
NUMBER OF SEQ ID NOS: 122
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ben-Sasson, Shmuel
TITLE OF INVENTION: SHORT PEPTIDES FROM THE A-REGION
TITLE OF INVENTION: PROTEIN KINASES WHICH SELECTIVELY
TITLE OF INVENTION: ACTIVITY
ELE REFERENCE: 1242.0003-000
CURRENT APPLICATION UNMBER: US/09/734,520
CURRENT FILING DATE: 2000-12-11
NUMBER OF SEQ ID NOS: 122
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 4; Conserv
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Query Match
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Fak
NAME/KEY: MYRISTATE
LOCATION: (1)...(0)
NAME/KEY: AMIDATION
LOCATION: (0)...(11)
                                                                                                                                                                                                                                                                                                         APPLICANT: Ben-Sasson, Shmuel
TITLE OF INVENTION: SHORT PEPTIDES FROM THE A-REGION OF
TITLE OF INVENTION: PROTEIN KINASES WHICH SELECTIVELY MODULATE PROTEIN KINASE
TITLE OF INVENTION: ACTIVITY
FILE REFERENCE: 1242.2003-000
                                                                                     OTHER INFORMATION: Ret NAME/KEY: MYRISTATE LOCATION: (1)...(0)
                                                                                                                                           LENGTH: 11
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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Pred. No.
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Pred. No. 1
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RESULT 15
US-10-012-034A-72
Sequence 72, Application US/10012034A
Patent No. US20020137141A1
GENERAL INFORMATION:
APPLICANT: Ben-Sasson, Shmuel
TITLE OF INVENTION: SHORT PEPTIDES FROM THE A-REGION OF
TITLE OF INVENTION: PROTEIN KINASES WHICH SELECTIVELY MODULATE PROTEIN KINASE
TITLE OF INVENTION: ACTIVITY
FILE REFERENCE: BEN-SASSON-5A
CURRENT APPLICATION NUMBER: US/10/012,034A
CURRENT APPLICATION NUMBER: 09/734,520
PRIOR APPLICATION NUMBER: 09/734,520
PRIOR APPLICATION STEE 2000-12-11
UNUMBER OF SEO ID NOS: 13
SOFTWARE: FRESURE OF WINDOWS Version 4.0
SEO ID NO 72
LENGTH: 11
TYPE: PRI
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Fak
Search completed: February 8, 2003, 11:02:27 Job time : 11 secs
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                                                                                                                                                                                                                      ; FEATURE:
; NAME/KEY: AMIDATION
; LOCATION: (0)...(11)
US-10-012-034A-72
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Best Local Similarity 80.0
Matches 4; Conservative
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7 HIVKL 11
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7 HVIKL 11
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1; Mismatches 0; Indels
                                                                                                                                               Score 24; DB 12; Length 11; Pred. No. 1.1e+02; Indels 1; Mismatches 0; Indels
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GenCore version 5.1.3 Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: February 8, 2003, 10:54:26; Search time 11 Seconds (without alignments) 41.476 Million cell updates/sec

Title: Perfect score: Sequence: US-09-685-010-30 54 1 KQKIKHVVKLK 11

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched:

Total number of hits satisfying chosen parameters:

411

112892 seqs, 41476328 residues

Minimum DB seq length: 0 Maximum DB seq length: 11

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_40:*

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

33000000000000000000000000000000000000		Result
10000000000000000000000000000000000000	18 17 15	Score
27.8 27.8 27.9 27.9 27.9 27.9 27.9 27.9 27.9 27.9	3. 7.	% Query Match
100 110 110 111 111 111 110 100 100 100	10 10	Length DB
AH3_PRUSE XYNB_DICB4 FARP_LOCMI FARP_LOCMI NEMS_DROME CSIS_BACSU Q20A_COMTE FAR2_ASCSU RS10_SERMA FARP_NANSE SYK_CAMUP TKNB_ONCMY UXA6_CHLTR HY7_PIG LPMS_TABE LY7_PIG LPMS_TABE MOSH_CLTAJA CXL1_COMMR UN05_CLOPA ACI_THUAL SPI_HALRO TKL4_LOCMI BPP7_BOTIN BPP7_BOTIN WWA2_ACHFU B44K_PORGI FUSS_FUSSO NEUX_HUMSN ULAK_MOUSE XYLA_STRSO COXM_RAT	LCMS_LEUMA COXK_ONCMY ULAD_HUMAN	B ID
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"Identification of tissue-specific isoforms for subunits Vb and VIIa

LCMS_LEUMA LCMS_LEUMA AC P21144; P414 DT O1-PEB-1994 DT O1-PEB-1994 DT O1-PEB-1994 DT O1-PEB-1994 DT O1-PEB-1994 DT O1-PEB-1994 DT O1-NOV-1995 DE Leucomyosupp OS Leucophaea m OC Eukaryota; pto OC Blaberoidea; Pto OC Blaberoidea; NCBLTTAXID-6 RN FIGURENCE, AN RP SEQUENCE, AN RP SEQUENCE, AN RP SEQUENCE, AN RP SEQUENCE OF PROTODEL KW MOD_RES FT MOD_RES FT MOD_RES SO SEQUENCE 1 Query Match Best Local Simi Matches 3; Matches 3; Matches 1; DO 1-OCT-1994 DT 01-OCT-1994 DT

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AH3_PRUSE
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T 01-DEC-1992 (Rel. 24, Last sequence update)
T 01-DEC-1992 (Rel. 35, Last annotation update)
T 01-NOV-1997 (Rel. 35, Last annotation update)
Maygdalin beta-glucosidase II (EC 3.2.1.117) (Amy 12 1502yme II) (AH II) (Fragment).
E 1502yme II) (AH II) (Fragment).
S Prunus serotina (Black cherry).
C Eukaryota; Viridiplantae; Streptophyta; Embryoph.
C Spermatophyta; Magnoliophyta; eudicotyledons; convested II; Rosales; Rosaceae; Amygdaloideae; Pr
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Eur. J. Biochen
-!- FUNCTION: 1
                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=94147969; PubMed=8313870;
Hughes G.J., Frutiger S., Paquet N., Pasquali C., Stughes G.J., Frutiger S., Paquet N., Postrasser Tissot J.-D., Bairoch A., Appel R.D., Hochstrasser "Human liver protein map: update 1993.";
Electrophoresis 14:1216-1222(1993).
-i- MISCELALANEOUS: ON THE 2D-GEL THE DETERMINED PI PROTEIN IS: 6, ITS MW IS: 15 kDa.
NCBI_TGID=23207;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Unknown protein from 2D-page of liver tissue
                                                                                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                  SWISS-2DPAGE;
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P31929;
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Oxidoreductase; Inner membrane;
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J. BIOCHEM. 221:1111-1116(1994).
FUNCTION: THIS PROTEIN IS ONE OF THE NUCLEAR-CODED POLYPEPTIDE CHAINS OF CYTOCHROME C OXIDASE, THE TERMINAL OXIDASE IN MITOCHONDRIAL ELECTRON TRANSPORT.
CATALYTIC ACTIVITY: 4 ferrocytochrome c + O(2) - 4 ferricytochic + 2 H(2)O.
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SIMILARITY: BELONGS TO THE CYTOCHROME C OXIDASE VII
                                                                                                                                                                                                                                 KKQTYHI 9
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2; Conserv
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1129 MW;
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1174 MW;
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Pred. No. 1.8e+03;
1; Mismatches 5;
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ledons; core eu
bideae; Prunus.
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                                                                                          (Amygdalin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (Spot 106) (Fragment).
                                                                                                                                                                                                                                                                                                              Length 9;
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r D.F.;
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RESULT 5

XYNB_DD

XYNB_DD

AC P80717

P80717

DT 01-NOV

DT 15-JUN

DE Endo-1

DE (1,4-b)

OS Dictyo

OC Bacter

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Best I
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Best Local :
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                  FARP_LOCMI STANDARD; PRT; 10 AA. P3855; 01-OCT-1994 (Rel. 30, Created) 01-OCT-1994 (Rel. 30, Last sequence update) 30-MAY-2000 (Rel. 39, Last annotation update) SchistoFLRFamide (PDVDHFLRF-amide) (Cardioexcitatory Locusta migratoria (Migratory locust), and Schistocerca gregaria (Desert locust).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P80717;
01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
15-JUN-2002 (Rel. 41, Last annotation update)
15-JUN-2003 (Rel. 41, Last annotation update)
15-JUN-2008 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=Seed;
Li C.P., Swa
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P80717;
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Adamsen A.K., Jacobsen S., Ahring B.K., Submitted (OCT-1996) to the SWISS-PROT-:- CATALYTIC ACTIVITY: Endohydrolysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (1,4-beta-D-xylan xylanohydrolase B) (Fragi
Dictyoglomus sp. (strain B4A).
Bacteria; Dictyoglomus group; Dictyoglomus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Xylan degradation; Hydrolase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR001000; Glyco_hydro_10.
PROSITE; PS00591; GLYCOSYL_HYDROL_F10; PARTIAL
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NON_TER
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"Prunus serotina amygdalin hydrolase a
Plant Physiol. 100:282-290(1992).
-i- CATALYTIC ACTIVITY: (R)-amygdalin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=69007;
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PATHWAY: Xylan degr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HYDROLASES)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   glucose.
SUBUNIT: MONOMER.
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                                                                                                                                                                                                                                                                                                                               KHVVKLK 11
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2; Conserv
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Metazoa;
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AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
E.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1033 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1144 MW;
Arthropoda;
                                                                                                                                                                                                                                                                                                                                                                                                   27.8%;
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40.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S., Ahring B.K.; the SWISS-PROT
                                                                                                                                                                                                                                                                                                                                                                        Score 15;
Pred. No. 4
2; Mismatch
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Pred.
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Mandibulata;
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No.
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of 1,4-beta-D-xylosidic
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4.1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                   .1e+03;
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Pancrustacea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CRC64;
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                                                               neuropeptide).
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NEMS_DROME
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Best I
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SEQUENCE.
SEQUENCE.
SPECIES-S.bullata; TISSUE-Head;
SPECIES-S.bullata; TISSUE-Head;
MEDLINE-93047886; PubMed=1358537;
Fonagy A., Schoofs L., Proost P.,
Fonagy A., Schoofs L. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SPECIES-S.gregaria; TISSUE-Thoracic nervous sy MEDLINE-89246543; PubMed-2719702; Robb S., Packman L.C., Evans P.D.; "Isolation, primary structure and bioactivity FMRF-amide-like neuropeptide from the locust,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  male
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de Loof
                                                                                                                                                                                  SPECIES=D.melanogaster;
MEDLINE=93002195; PubMe
                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapo
Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID-7227, 7385;
                                                                                                                                                                                                                                                                                                                                                               01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Dromyosuppressin (Neomyosuppressin) (NEB-MS).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neuropeptide;
MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE.
SPECIES=L.migratoria;
                                                                                                                                 TDVDHVFLRFamide
                                                                                                                                                                                                                                                                                                                    Drosophila melanogaster (Fruit Sarcophaga bullata (Grey flesh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gregaria."
Biochem, B
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                                                                                                                                                   "Isolation and
                                                                                                                                                                    Nichols
                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                       DMS OR NEMS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          NEMS_DROME
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Acridomorpha; Acridoidea; Acrididae; Oedipodinae; Locusta.
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                                                                                                                   MOL.
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TISSUE SPECIFICITY: FOUND
THE SALIVARY GLANDS, THE I
SIMILARITY: BELONGS TO THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   QF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       accessory glands,
uct.":
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A32543; A32543.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      m. Biophys. Res. Commun. 150:830-630. IN THE NEURAL CONTRO. MCTION: MUSCLE INHIBITING AGENT. INVOLVED IN THE NEURAL CONTRO. THE VISCERSAL MUSCLES OF THE HEART, ACCESSORY GLANDS AND TIDUCT. MAY BE INVOLVED IN THE REGULATION OF SALIVA SECRETION. ISSUE SPECIFICITY: FOUND IN AXONS OF THE MALE ACCESSORY GLANDS, THE HEART, AND THE OVIJUCT. HE SALIVARY GLANDS, THE HEART, AND THE OVIJUCT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
2; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14:409-421(1993)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AA;
                                                                                                                               structural characterization of Drosophila and FMRFamide-containing neural peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                         STANDARD;
                                                                                                              and FMRFamide-containing i. 3:213-218(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7010;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ria; TISSUE=Brain;
PubMed=7687352;
                                                                                                                                                                                  PubMed=1390001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1244 MW;
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50.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14; DB Pred. No. 6.3e 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and synthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AMIDATION.
D3C51729D2C1EAB2
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Neuropeptide; Amidation
MOD_RES
STRAIN=63;
MEDLINE=96035889; F
Schach S., Tshisuak
"Quinoline 2-oxidor
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SEQUENCE
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16-0CT-2001 (Rel. 40, Last annotation update)
Quinoline 2-oxidoreductase, alpha chain (EC 1.3.99.17) (Fragment).
Comamonas testosteroni (Pseudomonas testosteroni).
Bacteria; Proteobacteria; beta subdivision; Comamonadaceae; Co
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Cold shock protein CSI5 (11 kDa cold s
Bacillus subfilis.
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CBI_TaxID=1423;
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MEDLINE=96345629; PubMed=8:
Graumann P., Schroeder K.,
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SPECIES-A.suum;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CATALYTIC ACTIVITY: Quinoline + acceptor + 1(2H)-one + reduced acceptor. COFACTOR: FAD, MOLYBDENUM AND IRON-SULFUR. PATHWAY: FIRST STEP IN THE DEGRADATION OF COFACTOR: THE PATHWAY: FIRST STEP IN THE DEGRADATION OF COFACTOR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (3-METHYL-)QUINOLINE.
SUBUNIT: HETEROHEXAMER OF TWO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 noline and 3-methylquinoline degradation."
J. Blochem. 232:336-544(1995).
FUNCTION: CONVERTS (3-METHYL-)-QUINOLINE
1,2-DIHYDROQUINOLINE.
                                                                                                                                                                                                                                                                                                                                                                         FAMILY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VVKLK 11
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                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ascaris.
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                                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                             24.18;
                                                                                                                                                                                                                                                                                             MW.
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69D4073B5B11E350
                                                                                                                                                                                             0;
                                                                                                                                                                                                                 Score 13;
Pred. No.
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1; Mismatches
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thes 0;
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RESULT 12
FARP_MANSE
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Best Local S
Matches 2
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01-FEB-1994 (Rel. 28, Last sequence up
01-NOV-1995 (Rel. 32, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FARP_MANSE
P18523;
                                                                                                                                                                                                                                                                                                       Manduca sexta (Tobacco hawkmoth) (Tobacco hornworm). Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Insecta; Pterygota; Neoptera; Endopterygota; Lepidoptera; Ditrysia; Sphingiodea; Sphingidae; Sphinginae; Manduca. NCBI_TaxID=7130;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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30-MAY-2000
30-MAY-2000
15-JUN-2002
 Amidation;
                  PIR; A43977;
                                                       ÷
                                                                                                                            Peptides 11:849-856(1990)
                                                                                                                                                                                                 Hunt D.F.;
                                                                                                                                                                                                                MEDLINE-91045350; PubMed-2235684;
Kingan T.G., Teplow D.B., Phillips J.M.,
Hildebrand J.G., Homberg U., Kammer A.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ribosomal protein. NON_TER 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; AF058451; AAC14294.1; ...
Interpro; IPR001848; Ribosomal_S10,
PROSITE; PS00361; RIBOSOMAL_S10; PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation
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Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases
-!- FUNCTION: Involved in the binding of tRNA to the ri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Serratia.
NCBI_TaxID=615;
                                                                                                                                                             "A new peptide in the FMRFamide hawkmoth, Manduca sexta.";
                                                                                                                                                                                                                                                                         SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                  FMRFamide-like neuropeptide.
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                                   SIMILARITY: FAMILY.
                                                                      FUNCTION: INCREASES THE FORCE OF THE MAJOR POWER-PRODUCING FLIGHT MUSCLES AND SO IS LIKELY TO PLAY FLIGHT BEHAVIOR PATTERNS.
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SIMILARITY: BELONGS TO THE S10P FAMILY
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2; Conserv
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Neuropeptide
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                    A43977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                     BELONGS
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1214 MW;
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Last annotation
S10 (Fragment).
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                                                     THE
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Pred. No. 1.1e+05
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                                                     (FMRFAMIDE RELATED PEPTIDE)
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                                                                                     NEURALLY EVOKED CONTRACTIONS IN MUSCLES, THE DORSAL LONGITUDINAL A ROLE IN SUSTAINING OR PROMOTING
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, Jardine I.,
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                                                                                                                                                                                                                                                                                                                                                             Pancrustacea;
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Griffin
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RESULT 14
TKNB_ONCMY
ID TKNB_O
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DT 01-DEC
DT 01-DEC
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SYK_CAMP
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Best Local S
Matches 2
TKNB_ONCMY
P28500;
01-DEC-1992
01-DEC-1992
30-MAY-2000
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MOD_RES
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gene 183:219-224(1996).
-!- CATALYTIC ACTIVITY:
+ L-1ysyl-trna(Lys).
                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR002106; AAtRNA_ligaseII.
PROSITE; PS50862; AA_TRNA_LIGASE_II; PARTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; L77076; AAB41342.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entities requires a license agreement (
or send an email to license@isb-sib.ch)
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Bacteria; Proteobacteria;
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15-DEC-1998 (Rel.
15-JUN-2002 (Rel.
                                                                                                                                                                                                                                                                                                                                                             Aminoacyl-tRNA synthetase; Protein biosynthesis; Ligase; NON_TER 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -!- SUBUNIT: HOMODIMER (BY SIMILARITY).
-!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: BELONGS TO CLASS-II AMINOACYL-TRNA SYNTHETASE FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genome
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upsaliensis fur and its
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RESULT 15
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SEQUENCE
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01-0CT-1994 (Rel. 30, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Unknown protein from 2D-page from elementary body
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tachykinin; Neuropeptide; Amidation MOD_RES 10 10 AMIDAT
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"Substance-P-related and neurokinin-A-related of the cod and trout.";
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Actinopterygii; Neopterygii; Teleostei; Euteleo
Protacanthopterygii; Salmoniformes; Salmonidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Neurokinin A (Substance K) (Neuromedin L). Oncorhynchus mykiss (Rainbow trout) (Salmo Gadus morhua (Atlantic cod).
                                                                                                       Siena-2DPAGE; P38007;
                                                                                                                                        Submitted (SEP-1994) to the SWISS-PROT data
                                                                                                                                                           STRAIN=L2/434/Bu;
Bini L., Santucci A., Magi
Comanducci M., Christianen
                                                                                                                                                                                                                     Chlamydia trachomatis.
Bacteria; Chlamydiales; Chlamydiaceae; Chlamydia
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01-OCT-1994
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Interpro; IPR002040; Tachykinin.
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FUNCTION: TACHYKININS ARE ACTIVE PEPTIDES WHICH EXCITE EVOKE BEHAVIORAL RESPONSES, ARE POTENT VASODILATORS AND SECRETAGOGUES, AND CONTRACT (DIRECTLY OR INDIRECTLY) M
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Vtretou E., Ratti
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sp_bacteria:*
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09fec0 hordeum vul
09jn49 staphylococ
09r9c4 borrelia bu
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beet soil-	Q9dsn4 beet soil-b	Q9e8q3 beet soil-b	beet	1 beet soil-	Q9e8q5 beet soil-b	Q9dsn6 beet soil-b	Q9e8q6 beet soil-b	7 beet :	human	μ.	l mus m	7 trit:		Q9gd47 hydriastele		Q9n6m5 toxoplasma	Q9qyf6 mus musculu		077907 oreochromis	077897 oreochromis	oreoc	Q9h4h5 homo sapien	bacil	Q9rbv0 pseudomonas	xanthob	P89269 xestia c-ni	4 rattus	Q8rue7 zea mays (m

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RESULT
Q34909
ID Q3
AC Q3
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Matches 3
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SEQUENCE
Q34909 PRELIMINARY;
Q34909;
01-NOV-1996 (TrEMBLrel. 01,
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01-NOV-1996 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Membrane protein (1 is 3rd base in codon) (Fragment).
Transmissible gastroenteritis virus.
Viruses; SSRNA positive-strand viruses, no DNA stage; Nidovirales;
Coronaviridae; Coronavirus.
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Q66205;
                                                                                                                                                                                                                                                                                                                          Saccharomyces Cerevisiae.";
Mol. Microbiol. 2:89-99(1988).
                                                                                                                                                                                                                                                                                                                                                          Britton P., Carmenes R.S., Page K.W., Garwes D.J., Parra F.; "Sequence of the Nucleoprotein Gene from a Virulent British Field Isolate of Transmissible Gastroenteritis Virus and its Expression in
                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. STRAIN=FS772/70;
                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=88216185; PubMed=2835592;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=11149;
                                                                                                                               \mathbf{L}
                                                                                                                                                              2 QKIKHVV 8
                                                                                                                               EKLLHMV 7
                                                                                                                                                                                                                                                                                                          Y00542; CAA68606.1;
                                                                                                                                                                                                 Similarity 3; Conserv
                                                                                                                                                                                                                                                                       7 AA;
                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                         869 MW;
                                                                                                                                                                                                                 37.0%;
42.9%;
                                                                                                                                                                                                                                                                         72C691E727233B70 CRC64;
         Created)
                                                                                                                                                                                                   Score 20; DB 12;
Pred. No. 6.7e+05;
3; Mismatches 1
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056413;

01-NOV-1996 (TrEMBLrel. 01, 01)

01-NOV-1996 (TrEMBLrel. 01, 01)

01-DEC-2001 (TrEMBLrel. 19, 01)
                 P83158;
P83158;
01-DEC-2001
01-DEC-2001
01-DEC-2001
Photosystem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MCCracken A., Uhlenbusch I., Gellissen G.;
"Structure of the cloned Locusta migratoria mitochor
restriction mapping and sequence of its ND-1(URF-1)
Curr. Genet. 11:625-630(1987).
EMBL; X05286; CAA28905.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=88223478; PubMed=2836084; MEDLINE=88223478; PubMed=2836084; Gellissen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta; Pterygota; Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha; Acridodea; Acrididae; Oedipodinae; Locusta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-87318208;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Escherichia coli.
Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plasmid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=7004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytochrome b (Fragment)
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01-DEC-2001 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Locusta migratoria
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|:||
5 KLKH 8
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                                                                                                                                                                                                                                                                                                                                  Similarity
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                   (TrEMBLrel. 19, Created)
(TrEMBLrel. 19, Last sequence update)
(TrEMBLrel. 19, Last annotation update)
I iron-sulfur center (Photosystem I subunit VII)
                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                            PRELIMINARY;
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80.0%;
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(Fragment).
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Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1;
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                   Score 20;
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                                                                                                                                                                                                                                                                                                                                                                                              447E8354A05339C3 CRC64;
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6.7e+05;
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Q9FEC0;
01-MAR-2001
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Apte S.K.,
Submitted (
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01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
DNA for ORF'S from chromosome XV (Fragment).
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Hordeum vulgare (Barley).
Eukaryota; Viridiplantae;
                                                                                        Mla1uORF
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Saccharomycetales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COQ3 AND YOL096C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yeast 11:975-986(1995).
EMBL; X83121; CAA58183.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Saccharomyces cerevisiae (Baker's yeast).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anabaena sp. (strain L31).
Bacteria; Cyanobacteria; Nostocales; Nostocaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=4932;
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2 HTVKI 6
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SUBCELULAR LOCATION: CYTOPLASMIC.
SIMILARITY: THE IRON-SULFUR CENTERS
'BACTERIAL-TYPE' 4FE-4S FERREDOXINS.
erPro; IPR001450; 4Fe4S_FERREDOXIN; PART
SITE; PS00198; 4FE4S_FERREDOXIN; PART
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e S.K., Uhlemann E., Schmid R., Altendorf K.; mitted (OCT-2001) to the SWISS-PROT data bank FUNCTION: APOPROTEIN FOR THE IRON-SULFUR CEN
                                                                                                                                                                                                                                                                                                                                                                               IHNVVKI 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
4; Conserv
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3; Conserv
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                                                                                                                   (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
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57.1%;
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Streptophyta; Embryophyta; Tracheophyta;
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                                                                                                                Last sequence update)
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I; Iron-sulfur;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 19; DB 3;
Pred. No. 6.7e+05;
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t chromosome XV c
n reading frames.
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of ermC.
Dacueria; Spirochaetales; Spirochaetaceae; Borrelia.
NCBI_TaxID=139;
[1]
                                          ogroc4:
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Plasmid cp32-2, possible partition proteins (Fragmer Borrelia burgdorferi (Lyme disease spirochete).
Plasmid cp32-2.
Plasmid cp32-2.
Plasmid cp33-2.
Plasmid cp33-2.
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
Plasmid pE194 ermC leader region (Fragment).
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Narayanan C.S., Dubnau D.;
"Evidence for the translational are
studies and structural analysis w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zhou F., Kurth J., Wei F., Elliott C., Vale G., Yahiaoui N., Keller B., Somerville S., Wise R., Schulze-lefert P.; "Cell-autonomous Expression of Barley Mlal Confers Race-specific Resistance to the Powdery Mildew Fungus via a Rarl Independent Signaling Pathway.":
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteria; Firmicutes; Staphylococcus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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AY009939; AAG37357.1;
AY009938; AAG37355.1;
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03097; CAA26883.1; -.
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Q9XSL0;
01-NOV-1999
01-NOV-1999
01-DEC-2001
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                              Dinitrogenase beta-subunit (Fragment). Rhizobium sp. cowpea (strain IRc78). Bacteria; Proteobacteria; alpha subdiv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stevenson B., Casjens S., Rosa P.; 
"Evidence of past recombination events among the genes encoding 
Erp antigens of Borrelia burgdorferi."; 
Microbiology 144:1869-1879(1998).
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STRAIN-B31;
MEDLINE=98361033; PubMed=9695920;
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Anim. Genet. 32:19-26(2001).
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Cosenza G., Mariani P., Pastore N., Masina
"An allele associated with a non detectable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bovidae; Caprinae;
NCBI_TaxID=9925;
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Capra hircus (Goat).
              Rhizobiaceae; Rhizobium
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NCBI_TaxID=400;
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Q9S618;
Q1-MAY-2000
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Q1-MAY-2000
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01-MAR-2002
01-MAR-2002
Urbach E., Chisholm S.W.;
Urbach E., Chisholm S.W.;
"Genetic diversity in Prochlorococcus populations
sorted from the Sargasso Sea and Gulf Stream.";
Limnol. Oceanog. 43:1615-1630(1998).
                                                                                                                                                                                                                                                                                                                                                 Helicobacter pylori:
minimal age.";
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Prochlorococcus.
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Urbach E., Chisholm
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Q1-MAR-2002
Q1-MAR-2002
Q1-MAR-2002
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SEQUENCE FROM N.A.
MORE M.I., Kirsch F.P., Rathjen F.G.;
More M.I., Kirsch F.P., Rathjen F.G.;
"Targeted ablation of NrCAM and ankyrin-B resu-
"thres leading to cataract formation.";
"ihres leading to the EMBL/GenBank/DDBJ
                                                                   "A novel eyes absent proteingene 0:0-0(2002).
EMBL: AF455148: AAL57875.1;
NON_TER 10 10
                                                                                                                                             Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Q99MG3;
O1-JUN-2001 (TrEMBLrel. 17,
O1-JUN-2001 (TrEMBLrel. 17,
O1-JUN-2001 (TrEMBLrel. 17,
                                                                                                        Fee B.E.,
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Mammalia; Eutheria;
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Mus musculus (
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                                                                                                                                     NCBI_TaxID=9606;
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Pred. No. 1.4e+0
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l; Mismatches
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                                                                                                                                               Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Sciurognathi; Muridae; Murinae; Mus
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Maximum DB
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Listing first
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length: 11
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1 KOKIKHVVKLK 11
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ALIGNMENTS	- 654 25 - 654 27 - 654	S-09-424-292-7 -08-318-892B-30 S-09-685-010-60
	Sequence 60, Appl Sequence 31, Appl Sequence 31, Appl Sequence 11, Appl Sequence 29, Appl Sequence 29, Appl Sequence 4056, App Sequence 76, Appl Sequence 5662, Appl Sequence 25, Appl Sequence 27, Appl Sequence 27, Appl Sequence 28, Appl Sequence 29, Appl Sequence 31, Appl Sequence 39, Appl Sequence 31, Appl Sequence 39, Appl Sequence 39, Appl Sequence 5611, Appl Sequence 5611, Appl Sequence 5611, Appl Sequence 2911, Appl Sequence 2911, Appl Sequence 2911, Appl Sequence 11704, Appl Sequence 21, Appl Sequence 23, Appl Sequence 24, Appl Sequence 23, Appl Sequence 23, Appl Sequence 24, Appl Sequence 24, Appl Sequence 24, Appl Sequence 25, Appl Sequence 26, Appl Sequence 26, Appl Sequence 27, Appl Sequence 27, Appl Sequence 28, Appl Sequence 29,	equence 7, 1 quence 30, 1 equence 60,

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US-08-318-892-3

Sequence 3, Application US/08318892

GENERAL INFORMATION:

APPLICANT: Turley, Eva A

TITLE OF INVENTION: HYALDRONAN RECEPTOR (RHAMM = RECEPTOR TITLE OF INVENTION: HYALDRONAN RECEPTOR (RHAMM = RECEPTOR TITLE OF INVENTION: PEPTIDES

NUMBER OF SEQUENCES: 29

CORRESPONDENCE ADDRESS: ADDRESS: ADDRESSE: C/O FISH & NEAVE

STATE: 1251 Avenue of the Americas

CITY: New York

COUNTRY: USA

ZIT: 10020

COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Wordeperfect 6.1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/318,892

FILING DATE: 1-DEC-1994

CLASSIFICATION: 536

ATTORNEY/AGENT INFORMATION:

NAME: Pierri, Margaret A

REGISTRATION NUMBER: 30,709

REFERENCE/DOCKET NUMBER: SIM-8
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7 US-08-318-892-3 7 US-08-318-892B-3 15 US-09-169-077-1 18 US-09-424-292-4 20 US-09-685-010-30 23 US-09-978-309A-30

Sequence 3, Appli Sequence 1, Appli Sequence 1, Appli Sequence 4, Appli Sequence 30, Appl Sequence 30, Appl

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RESULT 3
US-09-169-077-1
; Sequence 1, Acplication US/09169077
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GENERAL INFORMATION:
APPLICANT: Turley
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                                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Pierri, Margaret A
REGISTRATION NUMBER: 307/
                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/318,892B
FILING DATE: 1.DEC-1994
CLASSIFICATION: 536
                                                                                                                                                                                                                                   TOPOLOGY: 11 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: TUTLEY, EV& A
TITLE OF INVENTION: HYALURONAN RECEPTOR (RHAMM - RECEPTOR FOR
TITLE OF INVENTION: HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (212) 596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: sir
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 1251 AV CITY: New York STATE: N.Y.
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Local Similarity 100.0%;
nes 11; Conservative (
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596-9090
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Pred. No. 0.0071;
Mismatches 0;
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US-09-685-010-30
, Sequence 30, Application US/09685010
; GENERAL INFORMATION:
; APPLICANT: Turley, Eva A.
; APPLICANT: Cruz, Tony F.
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SOFTWARE: PatentIn version 3.1
; SEQ ID NO 4
; LENGTH: 11
; TYPE: PRT
; ORGANISM: Artificial Sequence
; PEATURE:
; OTHER INFORMATION: Synthetic
US-09-424-292-4
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                                                         Query Match
Best Local Similarity
Matches 11; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4, Application US/09424292 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/424,292
CURRENT FILING DATE: 2000-07-06
PRIOR APPLICATION NUMBER: PCT/CA98/00448
PRIOR FILING DATE: 1998-05-11
                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: CA 2,205,771
PRIOR FILING DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 36
                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Improved Delivery of Disease Modifiers FILE REFERENCE: 205375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: GB 9607441.4
FILING DATE: 10-APR-1996
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: HUINDBER OF SEQUENCES: 5
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
1 KQKIKHVVKLK 11
                               1 KQKIKHVVKLK 11
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                                                                  Conservative
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                                                                                100.0%;
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                                                                            Score 54; DB 18;
Pred. No. 0.0071;
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Pred. No. 0.0071;
                                                                  Mismatches
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                                                                                               DB 18;
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RESULT 7
US-09-424-292-7
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                                                                                                                                                                                                                                                                                                                                                                                                             US-09-978-309A-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-978-309A-30
                                                                                                                     Sequence 7, Application US/09424292 GENERAL INFORMATION:
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SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 30

LENGTH: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 30, Application US/09978309A
APPLICANT: Turley, Eva A
TITLE OF INVENTION: Improved Delivery of Disease
FILE REFERENCE: 205375
CURRENT APPLICATION NUMBER: US/09/424,292
CURRENT FILING DATE: 2000-07-06
PRIOR APPLICATION NUMBER: PCT/CA98/00448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Pastrak, Aleksandra
APPLICANT: Turley, Eva A.
TITLE OF INVENTION: Compositions and Methods for Treating Cellular
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders
TITLE OF INVENTION: Hyaladherin and Hyaluronans
FILE REFERENCE: 033352-010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/541,522
PRIOR FILING DATE: 2000-04-03
PRIOR APPLICATION NUMBER: US 60/127,457
PRIOR FILING DATE: 1999-04-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 09/685,010 PRIOR FILING DATE: 2000-10-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/978,309A CURRENT FILING DATE: 2002-04-04
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SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATING TITLE OF INVENTION: CELLULAR RESPONSE TO INJURY AND OTHER PROLIFERATING CELL TITLE OF INVENTION: DISORDERS REGULATED BY HYALADHERIN AND HYALURONANS FILE REFERENCE: 910130.401C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Cruz,
                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Peptide composition that binds a hyalauronan
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
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LENGTH: 10
                                                 Best Local Similarity
Matches 9; Conserv
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Best Local :
                                                                           Query Match
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PRIOR APPLICATION NUMBER: CA:
PRIOR FILING DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 36
                                                                                                                                                                                             TELEFAX: (212) 596-909
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                      NAME: Pierri, Margaret A
REGISTRATION NUMBER: 30709
REFERENCE/DOCKET NUMBER: SI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: New York
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CITY: 1
                                                                                                                                                        TYPE:
                                                                                                                               STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE:
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KQKIKHVVK
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                                                                                                                                                        amino acid
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                                                                                                                                                                     9 amino acids
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                                                  Conservative
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                                                                                                                                linear
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                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
                                                                                                                 peptide
                                                                                                                                           single
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100.0%;
                                                              83.3%; Score 45; DB 7; L
100.0%; Pred. No. 4.2e+06;
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Pred. No.
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RESULT 9 US-09-685-010-60

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CURRENT APPLICATION NUMBER: US/09/978,309A
CURRENT FILING DATE: 2002-04-04
PRIOR APPLICATION NUMBER: US 09/685,010
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: US 09/541,522
PRIOR FILING DATE: 2000-04-03
PRIOR APPLICATION NUMBER: US 60/127,457
PRIOR APPLICATION NUMBER: US 60/127,457
PRIOR FILING DATE: 1999-04-01
NUMBER OF SEO ID NOS: 84
SOFTWARE: FASTSEQ for Windows Version 4.0
SEO ID NO 60
LENGTH: 9
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Best Local Similarity
Thes 9; Conserve
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US-09-978-309A-60
                                      Sequence 31, Application US/08318892B
GENERAL INFORMATION:
APPLICANT: Turley, Eva A
TITLE-OF INVENTION: HYALURONAN REC
TITLE OF INVENTION: HYALURONAN MEI
NUMBER OF SEQUENCES: 41
                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 9; Conserv
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CURRENT APPLICATION NUMBER: US/09/685,010
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 72
SOFTWARE: FASTSEQ FOR WINDOWS Version 4.0
SEQ ID NO 60
LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Pastrak, Aleksandra
APPLICANT: Turley, Eva A.
TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response to
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated by
TITLE OF INVENTION: Hyaladherin and Hyaluronans
FILE REFERENCE: 033352-010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Cruz, Tony
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapien
CORRESPONDENCE ADDRESS:
ADDRESSME: c/o FISH & NEAVE
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                                                                HYALURONAN RECEPTOR (RHAMM = RECEPTOR FOR HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                  83.3%; Score 45; DB 23; 100.0%; Pred. No. 4.2e+06;
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PRIOR APPLICATION NUMBER: PCT/CA98/00448
PRIOR FILLWG DATE: 1998-05-11
PRIOR FPLICATION NUMBER: CA 2,205,771
PRIOR FILLWG DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin version 3.1
SEQ ID NO 8
LENGTH: 10
TYPE: PRT
                                           Query Match
Best Local Similarity P
Matches 8; Conservat
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                                                                                                                          US-09-424-292-8
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US-09-424-292-8
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                                                                                                                                                                                                                                                                                                                           Sequence 8, Application US/09424292
GENERAL INFORMATION:
APPLICANT: TULTey, Eva A
TITLE OF INVENTION: Improved Delivery of Disease Modifiers
FILE REFERENCE: 205375
CURRENT APPLICATION NUMBER: US/09/424,292
CURRENT FILING DATE: 2000-07-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (212) 596-9090 INFORMATION FOR SEQ ID NO: 31:
                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                       OTHER INFORMATION: Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: sinc
TOPOLOGY: linear
MOLECULE TYPE: pepti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Pierri, Margaret A REGISTRATION NUMBER: 30709 REFERENCE/DOCKET NUMBER: SITELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Rele
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 1-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 10020
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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1 KOKIEHVVEL 10
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9; Conserva
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0, Version #1.30
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80.0%;
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100.0%; Pred. No.
tive 0; Mismatc
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                                                                          Score 41;
Pred. No. 1
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                                                             Mismatches
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                                                           Gaps
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RESULT 13 US-09-424-292-11

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US-08-474-654-29
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; OTHER INFORMATION: Synthetic
US-09-424-292-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 29, Application US/08474654 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/424,292
CURRENT FILING DATE: 2000-07-06
PRIOR APPLICATION NUMBER: PCT/CA98/00448
PRIOR FILING DATE: 1998-05-11
PRIOR APPLICATION NUMBER: CA 2,205,771
PRIOR FILING DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 36
                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA;
APPLICATION NUMBER: JP Hei-6-268723
FILING DATE: 01-NOV-1994
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Turley, Eva A
TITLE OF INVENTION: Improved Delivery of Disease Modifiers
FILE REFERENCE: 205375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
                                                                                                              HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Natori, Shunji
APPLICANT: Takahashi, Haruo
APPLICANT: Kobayashi, Yohei
APPLICANT: Hoshino, Masaharu
                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: NOVEL POLYPEPTIDE AND USE THEREOF NUMBER OF SEQUENCES: 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 21
CITY: N.W.
STATE: Was
                LOCATION: 9
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: sir
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                                                                            NAME/KEY:
                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/474,654
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                            ENGTH:
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20037-3202
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                                                                                                                                                                                                              9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3: Sughrue, Mion, Zinn, Macpeak & Seas
2100 Pennsylvania Avenue
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                                                                                                                                                    linear
                                                                        Modified-site
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NO
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              /note= "The carboxyl terminus
carbamoyl group."
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Search completed: February Job time: 133 secs
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; HYPOTHETICAL:
US-08-474-654-39
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                                                                                                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP He.
FILING DATE: 01-NOV-1994
INFORMATION FOR SEQ ID NO: 39
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 39, Applicat GENERAL INFORMATION:
                                                                                                                    Matches
                                                                                                                                             Query Match
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/474,654
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: NOVEL POLYPEPTIDE AND USE THEREOF NUMBER OF SEQUENCES: 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 210
CITY: N.W.
STATE: Wash
                                                                                                                                 Local
                                                                                                                                                                                                                    STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S.A. ZIP: 20037-3202
                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                          1 KLKVVVKLK 9
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7; Conserv
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Kobayashi, Yohei
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                                                                                                                   Conservative
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77.8%;
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77.8%;
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                                                                                                                 Score 31; DB 8;
Pred. No. 4.2e+06;
1; Mismatches 1
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Pred. No. 4.2e+06;
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GenCore version 5.1.3

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OM protein - protein search, using sw model

Run on: February 8, 2003, 10:58:17; Search time 18 Seconds (without alignments)

49.548 Million cell updates/sec

Title: US-09-685-010-30

Perfect score: 54
Sequence: 1 KQKIKHYVKLK 11

Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5

Searched: 405691 seqs, 81078759 residues
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Minimum DB seq length: 0
Maximum DB seq length: 11

Total number of hits satisfying chosen parameters:

123350

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

Pending_Patents_AA_New:*

1: /cgn2_6/ptodata/2/paa/PCT_NEW_COMB.pep:*

2: /cgn2_6/ptodata/2/paa/US06_NEW_COMB.pep:*

3: /cgn2_6/ptodata/2/paa/US07_NEW_COMB.pep:*

4: /cgn2_6/ptodata/2/paa/US09_NEW_COMB.pep:*

5: /cgn2_6/ptodata/2/paa/US09_NEW_COMB.pep:*

6: /cgn2_6/ptodata/2/paa/US00_NEW_COMB.pep:*

7: /cgn2_6/ptodata/2/paa/US00_NEW_COMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	. 80	7	o,	5	4	ω	2	1	Result No.
23	23	23	23	23	23	23	23	23	23	23	23	23	23	23	23	23	23	23	23	23	23	24	24	27	31	Score
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11	11	11	11	11	11	10	10	10	10	10	10	10	10	9	9	9	9	9	9	8	8	9	8	10	9	Length :
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30930,	2562			Sequence 9742, Ap		Sequence 33057, A		12718,			Sequence 25627, A	12718	Sequence 406, App	Sequence 40827, A	33056,		40827,		9809,	Sequence 19345, A	19345	e 284,	6	11,	Sequence 263, App	Description

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40.7	40.7	40.7	40.7	40.7	40.7	40.7	42.6	42.6	42.6	42.6	42.6	42.6	42.6	42.6	42.6	42.6	42.6	42.6
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US-09-641-528A-40879	US-09-641-528A-33147	US-09-641-528A-9824	US-09-641-528-40879	US-09-641-528-33147	US-09-641-528-9824	US-09-390-225B-28	US-09-641-528A-40663	US-09-641-528A-33058	US-09-641-528A-32779	US-09-641-528A-30930	US-09-641-528A-25628	US-09-641-528A-25429	US-09-641-528A-19346	US-09-641-528A-9742	US-09-641-528A-3545	US-09-641-528-40663	US-09-641-528-33058	US-09-641-528-32779
Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence .	Sequence	Sequence
40879, A	33147, A	9824, Ap	40879, A	33147, A	9824, Ap	28, Appl	40663, A	33058, A	32779, A	30930, A	25628, A	25429, A	19346, A	9742, Ap	3545, Ap	40663, A	33058, A	32/19, A

RESULT 2 US-10-083-259-11 US-10-083-259-11 Sequence 11, Application US/10083259 GENERAL INFORMATION: APPLICANT: Nielsen, Peter E. APPLICANT: Gowercman, Birgit Kjæeldgaard APPLICANT: Good, Liam APPLICANT: Good, Liam APPLICANT: Beck, Frederik APPLICANT: Beck, Frederik APPLICANT: Malik, Leila APPLICANT: Wissenbach, Margit TITLE OF INVENTION: Modified Peptide Nucleic Acid (PNA) Molecules FILE REFERENCE: PANT-0301 CURRENT APPLICATION NUMBER: US/10/083,259	Query Match 57.4%; Score 31; DB 6; Length 9; Best Local Similarity 62.5%; Pred. No. 3.3e+05; Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0; Qy 1 KOKIKHVV 8 Db 1 KYKLKHIV 8	-263 3, Application US/10133210 ORMATION: DeLisi, Charles Berzofsky, Jay Gulukota, Kamalakar Vaccaro, Dennis Weng, Zhiping Zhang, Chao Weng, Chao Weng, Chao Weng, Chao WENTION: COMPOSITIONS THEREOF ENCE: BU-035AX PLICATION UMBER: US/10/133,210 LING DATE: 2002-04-26 SEQ ID NOS: 281 Patentin Ver. 2.0 63 Artificial Sequence ORMATION: Description of Artificial Sequence: -263 57 44: Score 31: DB 6: Length
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RESULT 4

US-10-211-088-284

Sequence 284, Application US/10211088

GENERAL INFORMATION:
APPLICANT: Bright, Gary R.
APPLICANT: Premkumar, D. David
APPLICANT: Chen, Yih-Tai

TITLE OF INVENTION: Novel Fusion Proteins And Assays For Molecular Binding
FILE REFERENCE: 01-1022-US
CURRENT APPLICATION NUMBER: US/10/211,088

CURRENT APPLICATION NUMBER: 60/309,395
PRIOR APPLICATION NUMBER: 60/309,395
PRIOR APPLICATION NUMBER: 60/309,395
PRIOR FILING DATE: 2001-08-01
PRIOR FILING DATE: 2001-08-01
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APPLICANT: BOGOCH, SAMUEL
APPLICANT: BOGOCH, ELENO
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PRIOR APPLICATION NUMBER: 09/689,155
PRIOR FILING DATE: 2000-10-12
NUMBER OF SEQ ID NOS: 160
SOFTWARE: Patentin version 3.1
SEQ ID: 00 11
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Best Local :
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TITLE OF INVENTION: REPLIKIN PEPTIDES AND ANTIBODIES THEREFORE
FILE REFERENCE: 09425/46905
CURRENT APPLICATION NUMBER: US/10/189,437
CURRENT FILING DATE: 2002-07-08
CURRENT FILING DATE: 2002-07-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/278,761 PRIOR FILING DATE: 2001-03-27 NUMBER OF SEQ ID NOS: 729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 10/105,232
PRIOR FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: 09/984,057
PRIOR FILING DATE: 2001-10-26
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR FILING DATE: 2001-07-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 8
TYPE: PRT
                   PRIOR FILING DATE:
                                   PRIOR APPLICATION NUMBER: 60/341,589
NUMBER OF
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mes 6; Conserv
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SEQ ID NOS: 366
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71.48;
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Pred. No. 3
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31;
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RESULT 6
US-09-641-528A-19345
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Best Local Similarity
Thehes 4; Conserva
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CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51504
SOFTWARE: PBstSEQ for Windows Version 4.0
SEQ ID NO 19345
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Matches 4; Conservative
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                                                                                                  TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMFILE REFERENCE: 2060.0100001
                                                                                                                                                      APPLICANT:
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                                                                 CURRENT APPLICATION NUMBER: US/09/641,528A CURRENT FILING DATE: 2000-08-15
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                              PRIOR APPLICATION NUMBER: US 60/172,705 PRIOR FILING DATE: 1999-12-10
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                NUMBER OF SEQ ID NOS: 51504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Peptide Derived from Human Papillomavirus
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FastSEQ for Windows Version 4.0
                                                                                                                                                      Celis, Esteb
Grey, Howard
                                                                                                                                                                                       Chesnut, Robert
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57.1%;
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Pred. No.
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Pred. No. 3.3e+05;
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CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51004
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 9809
                                                                                                                                                                                                                                                                                Sequence 33056, Application US/09641528 GENERAL INFORMATION:
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TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS FILTE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS FILE REPERENCE: 2060.010001
CURRENT APPLICATION NUMBER: US/09/641,528
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEO ID NOS: 51504
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 33056
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ORGANISM: Artificial Sequence
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4; Conservative
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Chesnut, Robert
Celis, Esteban
Grey, Howard
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Grey, Howard
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Chesnut, Robert
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57.1%;
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Pred. No. 3.3e+05;
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; OTHER INFORMATION: Peptide Derived from Human Papillomavirus US-09-641-528-40827
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PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51504
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 40827
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Matches
SEQ ID NO 9809
LENGTH: 9
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APPLICANT:
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                              APPLICANT: Grey, Howard
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528A
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51504
SOFTMARE: FASTSEQ for Windows Version 4.0
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ORGANISM: Artificial Sequence
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US-09-641-528A-40827; Sequence 40827, Application US/09641528A; GENERAL INFORMATION:
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TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528A
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR PRILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51504
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 33056
NUMBER OF SEQ ID NOS: 51504
SOFTWARE:\PastSEQ for Windows Version 4.0
SEQ ID NO 40827
                                                 APPLICANT: Sette, Alessandro
APPLICANT: Sudney, John
APPLICANT: Southwood, Scott
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
APPLICANT: Chesnut, Robert
APPLICANT: Grey, Howard
TITLE OF INVENTION: USING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
TITLE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528A
CURRENT FILLNG DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
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Best Local :
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TYPE: PRT
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ORGANISM: Artificial Sequence
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Celis, Esteban
Grey, Howard
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Southwood, Scott
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Pred. No. 3.36
3; Mismatches
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Pred. No. 3.3e+05;
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RESULT 14
US-09-641-528-12718
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                                                    NUMBER OF SEQ ID NOS: 51504
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 12718
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PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51504
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 406
LENGTH: 10
TYPE: PRT
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APPLICANT:
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Best Local Similarity
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Best Local (
                                                                                                           CURRENT APPLICATION NUMBER: US/09/641,528
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
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APPLICANT: Grey, Howard
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528
CURRENT FILING DATE: 2000-08-15
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                   LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
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4; Conservative
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Chesnut, Robert
Celis, Esteban
Grey, Howard
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chesnut,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Southwood,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sidney,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     42.68;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 23; DB Pred. No. 3.3e 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 23; DB 5;
Pred. No. 1.7e+02;
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RESULT 15

Sequence 25627, Application US/09641528

Sequence 25627, Application US/09641528

GENERAL INFORMATION:
APPLICANT: Sette, Alessandro
APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Chesnut, Robert
APPLICANT: Chesnut, Robert
APPLICANT: Chesnut, Robert
APPLICANT: Chesnut, Robert
APPLICANT: Chesnut, Robert
Indepticanti Grey, Howard
Courter of INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS

FILE REFERENCE: 2060,0100001

CURRENT APPLICATION NUMBER: US/09/641,528

CURRENT APPLICATION NUMBER: US/09/641,528

CURRENT FILING DATE: 1909-12-10

NUMBER OF SEQ ID NOS: 51504

SOFTWARE: FastSED for Windows Version 4.0

SEQ ID NO 25627

LENGTH: 10
                                                                                                                                                                                                                                              ; TYPE: PRT; ORCANISM: Artificial Sequence; FEATURE: OTHER INFORMATION: Peptide Derived from Human Papillomavirus US-09-641-528-25627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; FEATURE. ( )
; OTHER INFORMATION: Peptide Derived from Human Papillomavirus US-09-641-528-12718
Search completed: February 8, 2003, 11:02:10 Job time : 19 secs
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                                                                                                                                                               Query Match 42.6
Best Local Similarity 44.4
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 42.6%;
Best Local Similarity 44.4%;
Matches 4; Conservative
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                                                                                1 QVLHVLKRK 9
                                                                                                                         3 KIKHVVKLK 11
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                                                                                                                                                                                   42.68;
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                                                                                                                                                               Score 23; DB 5; Length 10; Pred. No. 1.7e+02; 3; Mismatches 2; Indels
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Run on:
                                                                                                          OM protein - protein search, using sw model
February 8, 2003, 10:57:16; Search time 14 Seconds (without alignments) 23.118 Million cell updates/sec
                                                                                                                                                                                      GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
```

Title: Perfect score: Sequence: US-09-685-010-30 54 1 KQKIKHVVKLK 11

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched:

83753

262574 seqs, 29422922 residues

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0
Maximum DB seq length: 11

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

Issued_Patents_AA:*
1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	80	7	σ	5	4	ω	2	_	Result No.
20	20	20	20	20	20	20	20	20	20	21	21	21	21	22	22	22	22	22	22	23	23	23	23	23	24	25	Score
37.0	37.0	37.0	7.	37.0	7.	7.	7	37.0	7.	8	8	8	8	40.7	40.7	40.7	40.7	40.7	40.7	42.6	٠	42.6	42.6	42.6	44.4	46.3	Query Match Length
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US-08-621-259A-133	US-08-621-803-166	US-08-541-964-4	US-08-594-447-5	US-09-128-967-11	9	US-08-869-506-11	US-08-974-549A-122	US-08-612-302A-11	US-08-240-514-11	US-08-584-043A-5	US-08-584-043A-4	US-08-584-043A-3	US-07-924-054-6	US-09-115-737-151	US-08-465-325-151	US-08-456-112B-16	US-08-097-830E-16	US-09-644-600-65	US-09-644-600-34	US-09-115-737-94	25-	US-08-434-120-95	US-08-193-521-1	US-07-861-458C-139	US-09-019-346A-5	US-08-477-928A-47	ID
Sequence 133, App	166	4,	5,	11,	e 8, 7	11, 4	122,	11,	11,	დ ა	4	Sequence 3, Appli	e 6, Ap	151,	1,	16,	e 16,	e 65,	34,	Sequence 94, Appl	94,	e 95,	1, Ap	139	Sequence 5, Appli	Sequence 47, Appl	Description

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20 37.0 8 2 US-U8-06-06-36A-36 20 37.0 8 4 US-09-016-366A-36 20 37.0 8 4 US-09-288-061-2 20 37.0 8 4 US-09-288-061-2 20 37.0 8 4 US-09-217-352-166 20 37.0 9 2 US-08-621-803-161 20 37.0 9 2 US-08-621-803-161 20 37.0 9 2 US-08-621-803-161 20 37.0 9 2 US-08-621-259A-127 20 37.0 9 2 US-08-621-259A-128 20 37.0 9 4 US-09-217-352-161 20 37.0 9 4 US-09-217-352-161 20 37.0 9 4 US-09-217-352-161 20 37.0 9 4 US-09-217-352-161 20 37.0 9 5 PCT-US95-09262-127 20 37.0 10 2 US-08-621-803-156 20 37.0 10 2 US-08-621-803-157 20 37.0 10 2 US-08-621-803-157 20 37.0 10 2 US-08-621-803-153	37.0 8 2 US-08-065-047-19 37.0 8 2 US-09-016-366A-36 Sequence 37.0 8 4 US-09-218-061-2 Sequence 37.0 8 4 US-09-217-352-166 Sequence 37.0 8 5 PCT-US95-09262-133 Sequence 37.0 9 2 US-08-621-803-160 Sequence 37.0 9 2 US-08-621-803-151 Sequence 37.0 9 2 US-08-621-803-151 Sequence 37.0 9 2 US-08-621-259A-127 Sequence 37.0 9 2 US-08-621-259A-127 Sequence 37.0 9 2 US-08-621-259A-128 Sequence 37.0 9 4 US-09-217-352-161 Sequence 37.0 9 4 US-09-217-352-161 Sequence 37.0 9 4 US-09-217-352-161 Sequence 37.0 9 5 PCT-US95-09262-128 Sequence 37.0 10 2 US-08-621-803-157 Sequence 37.0 10 2 US-08-621-803-157 Sequence 37.0 10 2 US-08-621-803-153 Sequence 37.0 10 2 US-08-621-803-123 Sequence	45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	82
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	US-09-016-564/-19 US-09-016-366A-36 US-09-016-366A-36 US-09-218-061-2 US-09-218-061-2 US-09-217-352-166 US-09-217-352-163 US-08-621-803-160 US-08-621-803-160 US-08-621-803-195 US-08-621-803-195 US-08-621-803-195 US-08-621-803-195 US-08-621-803-195 US-08-621-803-127 US-08-621-352-161 US-09-217-352-161 US-09-217-352-161 US-09-217-352-161 US-09-217-352-162 US-09-217-352-163 US-09-217-352-164 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-09-217-352-165 US-08-621-803-157 US-08-621-803-157 US-08-621-803-157 US-08-621-803-153 US-08-621-803-15	10	10	10	9	9	9	9	9	9	9	9	9	9	8	8	8	8	α
US-U8-665-64/-19 US-09-218-36A-36 US-09-218-36A-3 US-09-217-352-166 PCT-US95-09262-133 US-08-621-803-161 US-08-621-803-161 US-08-621-803-161 US-08-621-259A-127 US-08-621-259A-128 US-09-217-352-161 US-09-217-352-161 US-09-217-352-162 US-09-217-352-163 US-09-217-352-163 US-09-217-352-163 US-09-217-352-156 US-09-217-352-156 US-09-217-352-156 US-09-217-352-156 US-08-621-803-156 US-08-621-803-156 US-08-621-803-156	sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence	N	N	N	σı	G	4	4	4	N	N	N	Ν	N	υ	4	4	2	٨
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19, 2, A 166, A 133, 160, 161, 195, 127, 128, 156, 157, 128, 156,		App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	Appli	App1	Appi

Qy 3 KIKHVV 8 Db 3 KIKHLV 8	Query Match 46.3%; Score 25; DB 4; Length 8 Best Local Similarity 66.7%; Pred. No. 2e+05; Matches 4; Conservative 2; Mismatches 0; Indel	; MOLECULE TYPE; peptide	ີດ໌າ	; TYPE: amino acid	; LENGTH: 8 amino acids	: SECTION FOR SECTIONS: 4/:	; TELEFAX: (202) 639 7890	; TELEPHONE: (202) 639 7700	TELECOMMUNICATION INFORMATION:	REFERENCE/DOCKET NUMBER: 19060-0105	7n	~	; CLASSIFICATION: 536	; FILING DATE: 07-JUN-1995	CORRENT APPLICATION UNIMPER: IIS/08/477.9384	; SOFTWARE: PatentIn Release #1.0, Version #1.30	SYSTEM: PC-DOS/MS-DOS	COMPUTER: IBM PC compatible	: MEDIUM TYPE: Floppy disk	ZIP: 20004-2400	; COUNTRY: U.S.A.	 CTTY: Washington D C	IDOS DODONISTIS	S	R OF SEQUENCES: 49	INVENTION: LYMPHOCYTE MEDIATED IMMUNE	: TITLE OF INVENTION: METHODS FOR CONTROLLING T	RMATION:	; Patent No. 6207389	RESULT 1 US-08-477-928A-47	
	Length 8;); Indels															.30										RESPONSES					
	0;																									V 2					
	Gaps																														

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: SEQ ID NO 5
: LENGTH: 10
: TYPE: PRT
: ORGANISM: HIV-1 matr1x protein
US-09-019-346A-5
                                             TGSQLOGY:
US-07-861-458C-139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 3
US-07-861-458C-139
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Query Match_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 5, Application US/09019346A Patent No. 6372720 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION OF THE STORY APPLICATION UNBER: US709/019,346A
CURRENT FILING DATE: 1998-02-05
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Paten***-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 6232061
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 139, Application US/07861458C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Longmuir, Kenneth J.
APPLICANT: Waring, Alan J.
APPLICANT: Haynes, Sherry M.
TITLE OF INVENTION: Liposome Fure Reference: A65189/RFT/DAV
                                                                                                                                  TELEFAX: (617) 542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 04:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5970
                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM PS/2 Model 50Z or 55SX
OPERATING SYSTEM: MS-DOS (Version 5.0)
                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/861,458C
FILLING DATE: 04/01/92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Marchionni, Mark Andrew APPLICANT: Johnson, Carl D. TITLE OF INVENTION: HOWOLOGY CLONING NUMBER OF SEQUENCES: 142
CORRESPONDENCE ADDRESS: ADDRESSEE: Fish & Richardson STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A. ZIP: 02110-2804
                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Boston
                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 KQKIKH 6
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                                                                             amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    44.48;
       42.68;
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Pred. No. 1.8e+02;
         Score 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
         DB 4;
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         Length 7;
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US-08-193-521-1
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                                                         Matches
                                                                      Query Match
Best Local
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Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: Olstein, Elliot M.
REGISTRATION NUMBER: 24.02
REFERENCE/DOCKET NUMBER: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Maloy, W.
APPLICANT: Kari, U. I
APPLICANT: Williams,
                                                                                                                                                                                                                                                                     TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inc
                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
TELEFAX: 201-994-1744
                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Biologically Active Peptide TITLE OF INVENTION: Compositions and Uses There NUMBER OF SEQUENCES: 19
                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                            MOLECULE TYPE: peptide
                                                                                                                                 OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PS,
OPERATING SYSTEM:
                                                                                                                                                                                                           STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 07/7 FILING DATE: 13-SEP-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Roseland
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                       LENGTH:
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2 KKLKKLLKL 10
                            2 QKIKHVVKL 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4 IKHVV 8
                                                         4.
                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07068
                                                                                                                                                                                                                                        amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EE: Carella, Byrne, Bain, Gilfillan,
EE: Cecchi & Stewart
6 Becker Farm Road
                                                                                                                                                                                                                                                      11 amino acids
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                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Maloy, W. Lee
Kari, U. Prasad
                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inch diskette
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                                                                         42.6%;
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                                                                                                                                 May be a C-terminal amide, and/or may be acetylated at N-terminus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       US/07/870,960
                                                                                                                                                                                                                                                                                                                                                           24,025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 2e+05;
1; Mismatches
                                                         Score 23; DB 1; Length 11; 
pred. No. 2.9e+02; 
4; Mismatches 1; Indels
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                                                         Gaps
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RESULT 5 US-08-434-120-95 ; Sequence 95, Application US/08434120

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                                                                                                                                                                                         Sequence 94, Application US/08465325 Patent No. 5686563
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                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Jacob, Leonard S. APPLICANT: Maloy, W. Lee TITLE OF INVENTION: Treatment TITLE OF INVENTION: Malignanci TITLE OF INVENTION: Biological NUMBER OF SEQUENCES: 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 201-994-1744 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Olstein, Elliot M.
REGISTRATION NUMBER: 24,025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IBM PS/2
                                                  TITLE OF INVENTION: BITTLE OF INVENTION: N-'
NUMBER OF SEQUENCES: 1:
CORRESPONDENCE ADDRESS:
                                                                                                                 APPLICANT: Magainin Pharmaceuticals Inc.
APPLICANT: 5110 Campus Drive
APPLICANT: Plymouth Meeting, PA 19462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: PC-DOS SOFTWARE: DW4.V2 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: 201-994-1700
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nes 4; Conserv
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TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/226,108 FILING DATE: APPLICATION NUMBER: US/07/937,462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                  ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & ADDRESSEE: Dunner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Carella, Byrne, Bain, Gilfillan, Cecchi & Stewart
Street, N.W. Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Malignancies with
Biologically Active Peptides
                                                                                    N-Terminal Substitutions
                                                                                                    Biologically Active Peptides Having
                                                                                                                                                                                                                                                                                                                                                                                 42.68;
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                                                                      153
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                                                                                                                                                                                                                                                                                                                                                           Score 23; DB 1; Le
Pred. No. 2.9e+02;
4; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                               Length 11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; MOLECULE TYPE: US-08-465-325-94
                                                                                                                                                                                                                                                                                                                                                                                               Patent No. 6348445
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 94,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 18-JAN-94
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 07/89
FILING DATE: 01-JUN-92
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC
OPERATING SYSTEM:
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                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Biologically Active Peptides With Reduced
Toxicity in Animals and a Method for Preparing
 PRIOR
                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: U.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 4; Conserv
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APPLICATION NUMBER: US/09/115,737
FILING DATE: 15-Jul-1998
CLASSIFICATION: CURNOWN>
& APPLICATION DATA:
                                                                                                                                                                                                                                STREET: 1300 I Street, N.W. Suite
                                                                                                                                                              COUNTRY: USA
ZIP: 20005-3315
                                                                                                                                                                                                 STATE: D.C
                                                                                                                                                                                                               CITY: Washington
                                                                                                                                                                                                                                                              ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
                                                                                                                                                                                                                                                                                                                                                                                                                                  Application US/09115737
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SYSTEM: PC-DOS/MS-DOS
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Michael McLane
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                                                                                                                                                                                                                                                                                                                                                                                   Prasad Kari
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 42.68;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07/891,201
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 23; DB 1;
Pred. No. 2.9e+02;
4; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05387.0021-03000
                                                                                 Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 11;
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Same

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APPLICANT: O'Brien, Timothy J.

APPLICANT: Tenimoto, Hirotoshi
TITLE OF INVENTION: TADG-15: An Extracellular Serine Protease
TITLE OF INVENTION: Overexpressed in Carcinomas
FILE REFERENCE: D6064CIP/D
CURRENT APPLICATION NUMBER: US/09/644,600
CURRENT FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: 09/421,213
PRIOR APPLICATION NUMBER: 09/421,213
PRIOR FILING DATE: 1999-10-20
PRIOR FILING DATE: 1999-10-20
PRIOR FILING DATE: 1998-02-20
RIOR FILING DATE: 1998-02-20
NUMBER OF SEQ ID NOS: 98
                                                                                   RESULT 9
US-09-644-600-65
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MOLECULE TYPE: peptide;
SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-115-737-94
                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: Residues 119-127 of the TADG-15 protein US-09-644-600-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 8
US-09-644-600-34
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                                  Sequence 65, Application US/09644600 Patent N 6451500 GENERAL INFORMATION:
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Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
APPLICANT: "Janimoto, Hirotoshi
                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (202) 408-4000
TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: 94:
                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
                                                                                                                                                              1 KVKDALKL 8
                                                                                                                                                                                             3 KIKHVVKL 10
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REGISTRATION NUMBER: 32,984
REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 07/891,201 FILING DATE: 01-JUN-92 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 08/465,330 FILING DATE: 05-JUN-1995 APPLICATION NUMBER: 08/184,462 FILING DATE: 18-JAN-94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 11 amino acids
                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                40.78;
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                                                                                                                                                                                                                                             Score 22; DB 4;
Pred No. 2e+05;
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Pred. No. 2
                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                  Indels
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; FEATURE:
; OTHER INFORMATION:
US-08-097-830E-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 16, Application US/08097830E
Patent No. 5652211
GENERAL INFORMATION:
APPLICANT: PORTO, MASSIMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 09/421,213
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: 09/027,337
PRIOR FILING DATE: 1998-02-20
NUMBER OF SEQ ID NOS: 98
SEQ ID NO 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
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                                                                                                                               TELEFAX: (212) 302-89: INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: TADG-15: An Extracellular Serine Protease TITLE OF INVENTION: Overexpressed in Carcinomas FILE REFERENCE: D6064CIP/D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/644,600 CURRENT FILING DATE: 2000-08-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                           REGISTRATION NUMBER: 25,669
REFERENCE/DOCKET NUMBER: 57
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 302-8989
TELEFAX: (212) 302-898
                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                  CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION: NAME: Costigan, James V.
                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Peptides For Neutralizing The TITLE OF INVENTION: Toxicity of Lipid A NUMBER OF SEQUENCES: 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Disl
COMPUTER: IBM PS,
OPERATING SYSTEM:
                                                    TOPOLOGY: circular
                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       40.7%;
Local Similarity 50.0%;
tes 4; Conservation
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                                                                          : 10 amino acids amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                     Word Perfect 5.1
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                 sulfide bond between Cys and Cys
                                                                                                                                                                                                                                                                                                                                                                                  US/08/097,830E
                                                                                                                                                                                                                         25,669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.50 inch, 1.44 Mb storage
                                                                                                                                                                                                           576-003
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Pred. No. 2e+05;
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Query Match Best Local Similarity

40.7%;

Score 22; DB 1; Pred. No. 3.9e+02;

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Query Match
Best Local Similarity
5; Conserve
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US-08-465-325-151
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US-08-456-112B-16
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                                                                                                                                                           Patent No. 5686563
GENERAL INFORMATION:
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APPLICANT: Porro, Massimo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212) 302-89: INFORMATION FOR SEQ ID NO:
                                             APPLICANT: Magainin Pharmaceuticals Inc.
APPLICANT: 5110 Campus Drive
APPLICANT: Plymouth Meeting, PA 19462
TITLE OF INVENTION: Biologically Active Peptides Having
TITLE OF INVENTION: N-Terminal Substitutions
NUMBER OF SEQUENCES: 153
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 302-8989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Costigan, James V. REGISTRATION NUMBER: 25,669 REFERENCE/DOCKET NUMBER: 57
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SOFTWARE: Word Perfect 5.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Hedman, Gibson & Costigan STREET: 1185 Avenue of the Americas
STREET:
               ADDRESSEE:
                               ADDRESSEE:
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1300 I.
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                Dunner
                            Finnegan, Henderson, Farabow, Garrett &
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Street, N.W. Suite 700
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55.6%;
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Pred. No. 3.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/184
FILING DATE: 18-JAN-94
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/891
FILING DATE: 01-JUN-92
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 32,984
REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
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MEDIUM TYPE: Floppy disk
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 514
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                                                                                                                                            ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.3
CURRENT APPLICATION DATA:
             APPLICATION NUMBER: US/09/115,737
FILING DATE: 15-Jul-1998
CLASSIFICATION: -Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               D.C
                                                                                                                                                                                                                STATE: D.C
                                                                                                                                                                                                                                               Dunner, L.L.P.
STREET: 1300 I Street, N.W. Suite 700
                                                                                                                                                                                                 COUNTRY: USA
                                                                                                                                                                                                                                CITY: Washington
                                                                                                                                                                                                                                                                                 ADDRESSEE: Finnegan, Henderson, Farabow,
APPLICATION NUMBER: 08/465,330
                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/09115737
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                                                                                                                                                                                                                                                                                                                               Biologically Active Peptides With Reduced Toxicity in Animals and a Method for Preparing Same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05387.0021-03000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 22; DB 1;
Pred. No. 4.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 11;
                                                                                                                                                                                                                                                                                 Garrett &
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; SEQUENCE TYPE: peptide ; SEQUENCE DESCRIPTION: SEQ ID NO: 151: US-09-115-737-151
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US-07-924-054-6
                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/924,054
FILING DATE: 1920903
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: RESNICK, David S
NAME: RESNICK, David S
NAME: RESNICK, DAVId S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 6, Application US/07924054 Patent No. 5486472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                       REFERENCE/DOCKET NUMBER: 40
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)523-3400
TELEFAX: (617)523-6440
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: FORGLS, Jean B
REGISTRATION NUMBER: 32,984
REFERENCE/DOCKET NUMBER: 05387.0021-06000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4400
TELEPHONE: (202) 408-4400
INFORMATION FOR SEQ ID NO: 151:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: SUZUKI, NO. 5486472uhiro
APPLICANT: KITADA, Chieko
APPLICANT: TSUDA, Masao
TITLE OF INVENTION: ANTIBODY TO PACAP AND USE THEREOF
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: DAVID G. CONLIN; DIKE, BRONSTEIN, ROBERTS&
ADDRESSEE: CUSHMAN
-924-054.5
                    MOLECULE TYPE:
                                                                                                    SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U
ZIP: 02109
                                            TOPOLOGY:
                                                                               LENGTH:
                                                                                                                                                                                                                                             REGISTRATION NUMBER: 34235
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                                                            AMINO ACID
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TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 05-JUN-1995
APPLICATION NUMBER: 08/184,462
FILING DATE: 18-JAN-94
APPLICATION NUMBER: 07/891,201
FILING DATE: 01-JUN-92
                                                                           8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SD
                       peptide
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Pred. No. 4.3e+02;
4; Mismatches 2;
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                                                                                                                                                                                  TELEPHONE: (213) 489-16
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                        Matches
                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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Best Local Similarity
Matches 3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: January 8, CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Smith, Louis C.
APPLICANT: Sparrow, James APPLICANT: Hauer, Jochen
APPLICANT: Mims, Martha P.
                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
                                               38.98;
Local Similarity 55.68;
es 5; Conservation
                                                                                                                                         STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                    TYPE: amino acids
                                                                                                                                                                                                                                                                                      NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327 REFERENCE/DOCKET NUMBER: 21
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/584,043A FILING DATE: January 8, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 6.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                FILING DATE
                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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                          1 KQKIKHVVK 9
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California
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                                                                                                                          peptide
                                                                                                                                                       single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LIPOPHILIC PEPTIDES FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         James T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             38.9%;
                                                                    Score 21; DB 4;
Pred. No. 2e+05;
                                                                                                                                                                                                                                                                                       217/189
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Pred. No.
                                                     Mismatches
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2e+05;
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                                                                                  Length 9;
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                                                     Indels
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                                                     0;
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Search completed: February Job time : 15 secs

8,

2003, 10:59:26

0

GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: Pebruary 8, 2003, 11:02:17; Search time 14 Seconds (without alignments) 68.667 Million cell updates/sec

Title:
Perfect score:
Sequence: US-09-685-010-31 46 1 KLKSQLVKRK 10

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 283224 seqs, 96134422 residues

Total number of hits satisfying chosen parameters: 1100

Minimum DB seq length: 0 Maximum DB seq length: 10

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

pIR_73:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	80	7	6	v	4.	w	2	1	! !	Result		
12	12	12	12	12	12	12	12	13	13	13	13	13	13	13	13	13	13	13	13	14	14	14	14	14	15	15	15	16	Score			
26.1	26.1	•	•					•		28.3				٠	•	•	٠		•	•	•	30.4	30.4	30.4	•	32.6	•	•	. .	Query	æ	
9	9	9	œ	7	7	6	5	10	10	10	10	10	9	9	80	7	8	6	6	10	10	9	9	9	10	10	9	4.	Length			
N	N	N	N	N	N	N	N	N	N	N	N	ผ	N	N	N	N	N	N	N	N	N	N	N	۲	N	N	N	N	B			
PC7076	S78762	813333	I64832	PS0254	S57274	S02617	T14910	PX0060	S71948	C39111	S09387	D37397	S10920	A60108	PL0162	S55548	I48126	A37765	S11024	PQ0784	S28055	S15850	A60957	YFPG	C54226	A42089	I54379					SUMMARIES
	ribosomal protein	alpha/beta-gliadin	Ca2+-transporting	18K protein 5507 -	triacylglycerol li	alcohol dehydrogen	hypothetical prote	lysosome-associate	matrix metalloprot	leavy chair		hypothetical prote	venom protein HR-3	exotoxin A - Strep	ä	mcrB protein - Esc	=		hydrogensulfite re	NADH2 dehydrogenas	cytochrome b559 co	36-mon	te growth	ש	light-harvesting p	ъ.	Ę	phospholipase C (E	ption			500

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30
11	11	11	11	11	11	11	11	11	11	11	11	12	12	12	12
23.9	23.9	23.9	23.9	23.9	23.9	23.9	23.9	23.9	23.9	23.9	23.9	26.1	26.1	26.1	26.1
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N	N	N	N	N	N	۲	N	N	N	N	N	N	N	N	N
PC2197	S13889	B24362	S30494	S45651	PN0043	LFSAME	A38671	S19630	B33932	JH0784	A61419	S70721	S13224	S27178	I52974
zymogen granule me	phosphoenolpyruvat	chloramphenicol O-	cat gene leader pe	probable Na+-trans	phosphatidylethano	probable msrA lead	peptidylglycine mo	ribosomal protein	Ig mu chain D regi	neuropeptide TE-6	sarcosine dehydrog	heat shock protein	virG protein - Agr	neurokinin A-relat	seminal vesicle pr

DB 2; Length 9; 2.8e+05; ches 0; Indels	C;Species: Homo sapiens (man) C;Date: 01-Nov-1996 #sequence_revision 01-Nov-1996 #text_change 21-Jul-2000 C;Accession: 154379 R;Arai, E; Ikeuchi, T; Nakamura, Y. Hum. Mol. Genet. 3, 937-939, 1994 A;Title: Characterization of the translocation breakpoint on chromosome 22q12.2 in a patil A;Reference number: 154379; MUID:95038750; PMID:7951241 A;Accession: 154379 A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-9 <res> A;Cross-references: GB:S75841; NID:g861532; PIDN:AAD14190.1; PID:g4261890 A;Cross-references: GB:S75841; NID:g861532; PIDN:AAD14190.1; PID:g4261890</res>	ocal Similarity 75.0%; Pred. No. 2.8e+05; 3; Conservative 1; Mismatches 07 VKRK 10 : MKRK 4	A;Title: Role of the upstream region containing an intrinsic DNA curvature in the negative A;Reference number: 140870; MUID:92396045; PMID:1522810 A;Accession: I40870 A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-4 <res> A;Residues: 1-4 <res> A;Cross-references: EMBL:X62825; NID:g40622; PIDN:CAA44636.1; PID:g4377417 C;Genetics: A;Gene: plc C;Keywords: phosphoric diester hydrolase Query Match 34.8%; Score 16; DB 2; Length 4;</res></res>	
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C;Accession: C54226

R;Kerfeld, C.A.; Yeates, T.O.; Thornber, J.P.
Biochemistry 33, 2178-2184, 1994

A;Title: Purification and characterization of the peripheral antenna A;Reference number: A54226; MUID:94162224; PMID:8117674

A;Accession: C54226
                                 R;Pleau, J.M.; Dardenne, M.; Blouquit, Y.; Bach, J.F. J. Biol. Chem. 252, 8045-8047, 1977
A;Title: Structural study of circulating thymic factor: A;Reference number: A01523; MUID:78026571; PMID:914862
A;Accession: A01523
                                                                                                                                                                                      N; Alternate names:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        light-harvesting protein B-830 alpha-3 chain - Chromatium purpuratum (fragm
C;Species: Chromatium purpuratum
C;Date: 05-Jan-1996 #sequence_revision 05-Jan-1996 #text_change 05-Jan-1996
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A;Note: sequence extracted from NCBI backbone (NCBIN:82267,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Reference number: A42089; A; Accession: A42089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transcription factor I-POU protein, alternative splice form - fruit fly (Dr
C;Species: Drosophila melanogaster
C;Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 20-Mar-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         밁
 A; Molecule type: protein
A; Residues: 4-9 < PLE>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R, Treacy,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
                                                                                                                              ;Alternate names: FTS (facteur thymique serique);Species: Sus scrofa domestica (domestic pig);Date: 13-Jul-1981 #sequence_revision 13-Jul-1981 #text_change 07-May-1999;Date: 13-Jul-1981 #text_change 07-May-1999;Accession: A01523; A60983
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Keywords: antenna complex; light-harvesting polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                           l Similarity
3; Conserv
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3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a two amino acid difference in the I-POU homeodomain 089; MUID:92154665; PMID:1346754
                                                                                                                                                                                                                                                                                                                                                                                            32.6%;
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Pred. No. 7.6e+03;
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J. 7.6e+03;
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A;Gene:
C;Superf
C;Keywor
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                                                                                                                                                                                                                                                                                                                      RESULT
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C; Accession: $15850
C; Accession: $15850
R; Bergman, T.; Postlind, H.
Biochem. J. 276, 427-432, 1991
Biochem. J. 276, 427-432, 1991
A; Title: Characterization of mitochondrial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: protein
A;Molecule type: protein
A;Residues: 1-9 < ERN>
C;Comment: This peptide induces DNA synthesis in immature thymocytes, but not peripheral in a variety of immunoassays.
C;Comment: This peptide was isolated in two forms. One form contained the pyrrolidone can r form (thymocyte growth peptide) contains a large, non-peptide blocking group with a hig C;Superfamily: thymic factor
C;Keywords: blocked amino end; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Glx) (in FTS) #status experimental
F;1/Modified site: blocked amino end (Glx) (in thymocyte growth peptide) #status experime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R;Ernstroem, U.; Gafvelin, G.; Rudja, J.M.
Biosci. Rep. 10, 403-412, 1990
A;Title: Purification fthymocyte growth peptide (TGP)
A;Reference number: A60957; MUID:91064427; PMID:2249004
A;Accession: A60957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              thymocyte growth peptide - sheep
N;Contains: FTS (facteur thymique serique)
C;Species: Ovis orientalls aries, Ovis amm
C;Date: 31-Dec-1993 #sequence_revision 31-
C;Accession: A60957
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A;Residues: 'Z',2-4,'Z',6-9 <BAC>
C;Comment: This peptide induces DNA synthesis in
in a variety of immunoassays.
C;Comment: See PIR:A60957 (sheep) for discussion
C;Superfamily: thymic factor
C;Keywords: pyroglutamic acid
F;1/Modified site: pyrrolldone carboxylic acid (G
                                                                    A; Molecule type:
A; Residues: 1-9 •
C; Genetics:
                                                                                                                                                                                                                                                                     vitamin D3 26-monooxygenase (EC 1.14.14.-) cytochrome P450 27-
N;Alternate names: cytochrome P450(26); vitamin D3 26-hydroxyla:
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 23-Apr.1993 #sequence_revision 23-Apr-1993 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Bach, J.F.; Dardenne, M.; Pleau, J.M.; Rosa, J. Nature 266, 55-57, 197.
A;Title: Biochemical characterisation of a serum thymic A;Reference number: A60983; MUID:77123829; PMID:300146 A;Accession: A60983
                                                                                                                                      A; Reference number: S15850; A; Accession: S15850
Superfamily: unassigned cytochrome P450; cytochrome P450 homology; Keywords: heme; mitochondrion; monooxygenase; oxidoreductase; tra
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3; Conserv
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site: pyrrolidone
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                                                                                                                 protein
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Pred. No.
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31-Dec-1993
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o. 2.8e+05;
o;
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0. 2.8e+05;
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PMID: 2049072
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RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NADH2 dehydrogenase (EC 1.6.99.3) 29K chain - fava bean mitochondrion (fragn N;Alternate names: complex I 29K chain; NADH-ubiquinone reductase 29K chain C;Species: mitochondrion Vicia faba (fava bean) C;Date: 03-May-1994 #sequence_revision 07-Oct-1994 #text_change 03-Jun-2002 C;Accession: PQ0784
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                                                                                                                                                                                                                                                                                                             A;Molecule type: protein
A;Residues: 1-10 <LET>
C;Comment: Complex I, mitochondrial NADH-ubiquiquinone reductase, ranging from 5K to 75K.
                                                                                                                                                                                                                                                                                                                                                                                   A; Reference number: A; Accession: PQ0784
                                                                                                                                                                                                                                                                                                                                                                                                 A; Title: Purification and particles A; Reference number: PQ0775;
                                                                                                                                                                                                                                                                                                                                                                                                                                    R; Leterme, S.; Bout: Plant Physiol. 102,
hydrogensulfite reductase (EC 1.8.99.3) N;Alternate names: bisulfite reductase;
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Best Local (
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Keywords: chloroplast; ]
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photosynthesis;
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50.0%;
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Pred. No. 2.8e
0; Mismatches
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Pred. No. 1.2e
2; Mismatches
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chain 1 - Desul
desulfofuscidin
                                                                                                                                                                                       No. 1.2e+04;
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2.8e+05;
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                Desulfovibrio thermophilus (fragment)
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A;Molecule type: DNA
A;Residues: 1-6 <RES>
A;Cross-references: GB:
C;Genetics:
A;Introns: 3/3
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C;Date: 19-Mar-1997 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998
C;Accession: S11024
R;Fauque, G.; Lino, A.R.; Czechowski, M.; Kang, L.; DerVartanian, D.V.; Moura, J.J.G.; Le Blochim. Blophys. Acta 1040, 112-118, 1990
A;Title: Purification and characterization of bisulfite reductase (desulfofuscidin) from A;Reference number: S11024; MUID:9035276; PMID:2165817
A;Residues: 1-6 <FRU>C;Keywords: 1-6 <FRU>C;Keywords: oxidoreductase
                                                                                                                                                                                                                                                                 alpha-tubulin - Chinese hamster (fragment)
(;Species: Cricetulus griseus (Chinese hamster)
C;Date: 04-Sep-1997 #sequence_revision 07-Nov-1997 #text_change
C;Accession: I48126
R;Elliott, E.M.; Sarangi, F.; Henderson, G.; Ling, V.
Can. J. Biochem. Cell Biol. 63, 511-518, 1985
A;Title: Cloning of 11 alpha-tubulin gene sequences from the gen
A;Reference number: I48126; MUID:86001952; PMID:2931165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Theroux, S.J.; Redlinger, T.E.; Fuller, R.C.; Robinson, S.J.
J. Bacteriol. 172, 4497-4504, 1990
J.Title: Gene encoding the 5.7-kilodalton chlorosome protein of A;Reference number: A37765, MUID:90330558; PMID:2376566
A;Accession: A37765
A;Status: preliminary
                                á
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A; Residues: 1-6 <THE>
A; Cross-references: G
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Pred. No. 2.8e+05
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exotoxin A - Streptococcus pyogenes (strain C203.S) (fragment)

N;Alternate names; blastogen A; scarlet fever toxin

C;Species: Streptococcus pyogenes

C;Date: 10-Nov-1992 #sequence_revision 10-Nov-1992 #text_change 07-Feb-1997

C;Accession: A60108

R;Schlievert, P.M.; Gray, E.D.

Infect. Immun. 57, 1865-1867, 1989

A;Title: Group A streptococcal pyrogenic exotoxin (scarlet fever toxin) type A and blast A;Recession: A60108

A;Recession: A60108

A;Recession: A60108

A;Residues: 1-9 <SCH>
                                                                                                  Query Match
Best Local Similarity
Wardhes 3; Conserve
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A;Title: Phosphorylation of paramyosin.
A;Reference number: PL0162; MUID:90107385; PMID:2532591
A;Accession: PL0162
Beer
Matches
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A;Note: the sequence is the phosphorylated tryptic peptide
C;Comment: This protein is thought to exist as a dimer of two subunits, termed alpha-pax om the carboxyl-terminal end of the molecule. Only alpha-paramyosin is phosphorylated by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   paramyosin - northern quahog (fragment)
C;Species: Mercenaria mercenaria (northern quahog)
C;Date: 31-Dec.1990 #sequence_revision 31-Dec-1990 #text_change 11-May-2000
C;Accession: PL0162
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F;5/Binding site: phosphate (Ser) (covalent) #status experimental
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EMBO J. 14, 2661-2669, 1995
A;Title: McrB: a prokaryotic protein specifically recognizing DNA containing modified A;Reference number: S5548; MUID:95300799; PMID:7781618
A;Accession: S55548
A;Molecule type: protein
A;Residues: 1-7 <KRU>
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Keywords: exotoxin
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FARS PANRE
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THYP PIG
PSBE CAPAN
TKUI UREUN
PARI ASCSU
MALE KLEPN
RSIO SERMA
TKUB RANRI
LPMS STABP
LPCA STABP
LPCA STABU
ULAK MOUSE
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Q46464 campylobact P31929 homo sapien P56923 rana tempor P82661 panagrellus P01255 sus scrofa Q03367 capsicum an p40751 urechis uni P31889 ascaris suu Q05564 klebsiella G8936 serratia ma P29135 rana ridibu P23211 staphylococ P3684 staphylococ P3684 staphylococ P3684 staphylococ P3684 staphylococ P3686 staphylococ P3686 staphylococ P3687 staphylococ P3688 pinus musculu P30091 homo sapien P81298 haemonchus P31874 panagrellus P36960 carnobacter P80709 carcinus ma P81886 porphyromon P33564 mycobacteri P80709 carcinus ba P81886 porphyromon P33564 mycobacteri P80709 carcinus ba P81886 porphyromon P3032 chunnus obe Q9przl oncorhynchu P8032 concorhynchu P8032 acholeplasm P38007 chlamydia t P25154 oryctolagus P41875 panagrellus P805423 melolontha P14595 tabanus atr
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용 성	SQ ST	X	38888888	388888	RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR	22228		RES		
1 1 - E	NON TER SEQUENCE Query Match Best Local Matches	EMBL; L77 InterPro; PROSITE; Aminoacyl	ntti see	Gene 18	SEQUENCE FROM N.A SEQUENCE 43954 MEDLINE-97149302; Bourke B., Rashidd "Characterization localization in a	1 ~ 6 ~ .	15-DEC-1998 15-DEC-1998 15-JUN-2002 Lysyl-tRNA (Fragment).	RESULT 1 SYK_CAMUP ID SYK_CA		8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8
LKSQLVKRK : : LKSELKEDK	ທຜິ	7770	swiss-pror sen the Sw suropean Bi by non-profiled and the cies required and an emai	TIGHT TO THE STATE OF THE STATE OF THE SUBUNIT: HON SUBCELLULAR SIMILARITY:	* 5 C D 5 B	obact ia; P obact axID=	-1998 -1998 -2002 -RNA ent).	CAMUP		ουουουουουο
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	9CEA46AB13372 Score 21; DB Pred. No. 1.9 ; Mismatches	in sel	y is copyright. It is produnstitute of Bioinformatics There institute. There institutions as long as atment is not removed. Ulicense agreement (See httplicense@isb-sib.ch).	lys: ARIT	0; H.L., Ch ter upsal		ed) sequence update) annotation update) 5.1.1.6) (Lysinet	PRT; 10	ALIGNMENTS	HTF2 PERAM HTF TENMO RPCH PANBO RS1 ERWCH BUK CLOPA FAR2 CALVO PAR3 CALVO HUTU KLEAE LITR PHYRO NEF HVIZ8 NEUK HUMAN SAMP_MUSCA
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TEMK_RANTE
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Best Local
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P56923;
30-MAY-2000
16-OCT-2001
16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                              Amphibian skin; Antibiotic; Amidation; Multigene family.

MOD_RES 10 10 AMIDATION.

SEQUENCE 10 AA; 1123 MW; 390549B337272457 CRC64;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          temporaria.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Barra D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Simmaco M., Mignogna
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Skin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Temporin
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unknown protein from Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation update)
Unknown protein from 2D-page of liver tissue
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                                                                                                                                                                                                               2 LKSQL 6
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SIMILARITY: BELONGS TO THE BREVININ/ESCULENTIN/GAEGURIN/RUGOSIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FUNCTION: HAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LVKKO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Biochem. 242:788-792(1996).
CTION: HAS ANTIBACTERIAL ACTIVITY AGAINST GRAM-POSITIVE
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Last annotation updat
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                                                                                                                                                                                                                                                                                                             Score 15; DB 1;
Pred. No. 3e+03;
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Pred. No. 1.1e+05;
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P01255;
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P82661;
16-OCT-2001
16-OCT-2001
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                 PIR; A01523; YFPG.
                                                                                                                                                                                                                                                                                                                                                                                               from pig serum. II. Amino acid sequence.";
J. Biol. Chem. 252:8045-8047(1977).
-i- MISCELLANEOUS: THE BIOLOGICAL SOURCE(S) AND PHYSIOLOGICAL ACTIVITIES OF THIS PEFTIDE HAVE NOT BEEN DETERMINED.
PIR; A01523; YFPG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE.
MEDLINE=78026571; PubMed=914862;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sus scrofa (Pig)
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MOD_RES 9 9
SEQUENCE 9 AA; 1077 MW;
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Submitted (JUL-2000) to
-!- FUNCTION: MYOACTIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pleau J.-M., Dardenne M., Blouquit Y., E "Structural study of circulating thymic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE, FUNCTION, Moffet C.L., Marks
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Panagrolaimoidea; Panagrolaimidae; Panagrellus.
NCBI_TaxID=6233;
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(Rel. 40, Last annotation update)
ike neuropeptide PF5 (AMRNALVRF-amide)
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ilarity 100.0%;
Conservative
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N.J., Halton D.W.,
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Pred. No. 1.1e+
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MEDLINE=9309270; PubMed=1463853;
Kuntz M., Camara B., Weil J.-H., Schantz R.;
"The psbL gene from bell pepper (Capstcum annuum): plastid editing also occurs in non-photosynthetic chromoplasts.";
Plant Mol. Biol. 20:1185-1188(1992).
-!- FUNCTION: THLS B-TYPE CYTOCHROME IS TIGHTLY ASSOCIATED REACTION CENTER OF PHOTOSYSTEM II AND POSSIBLY IS PART
                                                                                                                                                                                            MEDLINE=93236558; PubMed=8476410;
Ikeda T., Minakata H., Nomoto K., Kubota I., M
"Two novel tachykinin-related neuropeptides in
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01-FEB-1995 (Rel. 31, Last sequence upon 101-NOV-1995 (Rel. 32, Last annotation
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PROSITE; PS00537; CYTOCHROME_B559;
Chloroplast; Photosystem II; Heme;
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                                                                                           Biochem. Biophys. Res. Commun. 19
-!- FUNCTION: CONTRACTILE ACTION
MUSCLE OF THE ANIMAL.
                                                                                                                                                   Urechis unicinctus.";
Biochem. Biophys. Res
                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Echiura; Xenopneusta; Urechidae; Urechis.
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Cytochrome B559 beta chain (Fragment).
  SEQUENCE
                              8
                                             Tachykinin; Neuropeptide; Amidation.
                                                                                                                                                                                                                                                                         TISSUE=Ventral
                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=6432;
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                                                                       -!- SIMILARITY: SOME SIMILARITY TO TACHYKININS
                                                                                                                                                                                                                                                                                                SEQUENCE, AND SYNTHESIS.
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SUBUNIT: DIMER OF AN ALPHA AND A BETA CHAIN.
SIMILARITY: BELONGS TO THE PSBE / PSBF FAMILY.
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     AA;
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Pred. No. 4.8e+0;
2; Mismatches
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817D0F59D6D69DC5
     C6D1C462C9D6C5A6 CRC64;
                                                                                                                     192:1-6(1993).
ON ON THE INNER CIRCULAR
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Electron
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Best Local S
Matches 3
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Best Local S
Matches 1
MEDLINE-93211295; PubMed-8459773;
MEDLINE-93211295; PubMed-8459773;
Machellier S., Perrin D., Hofnung M., Gilson E.;
Bachellier S., Perrin D., Hofnung M., Gilson E.;
"Bacterial interspersed mosaic elements (BIMEs) are present in the genome of Klebsiella.";
Mol. Microbiol. 7:537-544 (1993).

-i- FUNCTION: MALE IS INVOLVED IN THE HIGH-AFFINITY MALTOSE MEMBRAN TRANSPORT SYSTEM. INITIAL RECEPTOR FOR THE ACTIVE TRANSPORT OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KLEPN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
01-FBB-1996 (Rel. 33, Last annotation update)
FMRFamide-like neuropeptide AF1.
Ascaris suum (Pig roundworm) (Ascaris lumbricoides)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q05564;
01-OCT-1994
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MOD_RES 7 7
SEQUENCE 7 AA; 953 MW
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-i- TISSUE SPECIFICITY: FOUND IN THE NERVE CORDS AND A VARIETY OF GANGLIA PARTICULARLY IN THE ANTERIOR REGIONS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FAR1
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Bacteria;
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Maltose-binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                        Klebsiella.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Klebsiella
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1; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (Rel. 30, Created)
(Rel. 30, Last sequence update)
(Rel. 39, Last annotation update)
ding periplasmic protein (Maltodextrin-binding protein)
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69D40059CB144350 CRC64;
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Pred. No. 1.1e
4; Mismatches
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4.8e+03;
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                                                                                                                                                                                                                                                                                                                                Noorani S.M., Lindahl L., Zengel J.M.;
Submitted (APR-1998) to the EMBL/GenBank/DDBJ
-1- FUNCTION: Involved in the binding of tRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; X68329; CAA48406.1; -.
InterPro; IPR000567; SBP_BACTERIAL_1; PARTIAL.
                                                                                                                                             InterPro; IPR001848; Ribosomal_S10; PARTIAL.
                                                                                                                                                                      EMBL; AF058451; AAC14294.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                           Serratia marcescens
                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-MAY-2000
15-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Transport; Sugar transport; PerTplasmic.
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                       2 LKSQLVK 8
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                                                                                                                                                                                                                                                                                                       similarity).
SIMILARITY: BELONGS TO THE S10P FAMILY OF RIBOSOMAL PROTEINS.
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SUBCELLULAR LOCATION: Periplasmic.
SIMILARITY: BELONGS TO THE BACTERIAL EXTRACELLULAR SOLUTE-BINDING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROTEIN FAMILY 1.
MQNQRIR 7
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(Rel. 39,
(Rel. 41,
                                                    Conservative
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1214 MW;
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33.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                             gamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence update)
                                                   <u>υ</u>
                                                               Score 12;
Pred. No.
                                                                                                      DE3944004416D456 CRC64;
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Pred. No. 7.
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                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                              subdivision; Enterobacteriaceae;
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                                                                 1.1e+05;
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7.5e+03
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A to the rib
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RESULT 12
LPMS_STAEP
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TKNB_RANRI
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01-DEC-1992
16-OCT-2001
                                                                                                                                                                                                                                                                                                    LPMS_STAEP STANDARD; PRT; 8 AA
P23211;
01-NOV-1991 (Rel. 20, Created)
01-NOV-1991 (Rel. 20, Last sequence update)
01-MAR-1992 (Rel. 21, Last annotation updat
                                                                                                                  "Inducible erythromycin resistance in staphylococci is encoded member of the ATP-binding transport super-gene family."; mol. Microbiol. 4:1207-1214(1990).
                                                                                                                                                                    MEDLINE=91041730;
Ross J.I., Eady E.
                                                                                                                                                                                                                                                                           Probable msrA leader peptide. Staphylococcus epidermidis.
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InterPro; IPR002040; Tachykinin.
PROSITE; PS00267; TACHYKININ; 1.
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Burcher E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota, Metazoa, (
Amphibia; Batrachia,
NCBI_TaxID=8406;
                                                             This SWISS-PROT entry is copyright.
                                                                                                                                                                                                              SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rana ridibunda (Laughing frog) (Marsh
Eukaryota; Metazoa; Chordata; Craniata
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P29135;
                                                                                                                                                                                                                                    NCBI_TaxID=1282;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              related peptide from frog gut."; Biochem. J. 287:827-832(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Primary structure and receptor-binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=93075037; PubMed=1332683;
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        s SWISS-PROT entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the Ew European Bioinformatics Institute. There are no restr by non-profit institutions as long as its content if and this statement is not removed. Usage by and
                                                                                                     FUNCTION: MAY REGULATE EXPRESSION OF THE
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                                                                                                                                                         J.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                   Firmicutes;
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(Rel.
(Rel.
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24, Last seq
40, Last ann
                                                                                                                                                                    PubMed=2233255;
.A., Cove J.H.,
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AND CONTRACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hing frog) (Marsh frog).
Chordata; Craniata; Vertebrata;
Anura; Neobatrachia; Ranoidea;
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75.0%;
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annotation
agreement
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Pred. No. 1.2e
0; Mismatches
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noved. Usage by and for commercial
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Ranidae; Rana
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Matches 2
                        This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation—the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                  SPECIES=S.agalactiae; PLASMID=pIP501;
MEDLINE=9309867; PubMed=1461942;
Trieu-Cuot P., de Cespedes G., Horaud T.;
"Nucleotide sequence of the chloramphenicol resistance the streptococcal plasmid pIP501.";
                                                                                                                                                                                                                                                  Brueckner R., Matzura H.; "Regulation of the inducible chloramphenicol of the Staphylococcus aureus plasmid pUB112." EMBO J. 4:2295-2300(1985)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Schwarz S., Cardoso M.;
"Nucleotide sequence and pacetyltransferase encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Streptococcus agalactiae.
Plasmid pSC56, Plasmid pSC57, Plasmid pUB112, and Plasmid pIP501.
Bacteria; Firmicutes; Bacillales; Staphylococcus.
NCBI_TaxID=1280, 1311;
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01-OCT-1994
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                                                                                                                                                                                                                                                                                                                                                                                               "Nucleotide sequence and structural relationships of a chloramphenicol acetyltransferase encoded by the plasmid pSCS6
                                                                                                                                                                                                                                                                                                                                                                                                                                        SPECIES=S.aureus; PLASMID=pSCS6; MEDLINE=92388047; PubMed=1517170;
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MEDLINE=92027652; PubMed=1929326;
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Leader peptic
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  EMBL; M58515; AAA26612.1;
                                                                                                                                            Plasmid
                                                                                                                                                                                                                                                                                                           SPECIES=S.aureus; PLASMID=pUB112; MEDLINE=86081739; PubMed=3865770;
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CE 8 AA; 937 MW;
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L; Mismatches
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Best Local :
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15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
16-JUL-1998 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
Probable ribulose bisphosphate carboxylase/oxygenase activase
Probable vibulose bisphosphate carboxylase/oxygenase activase
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15-DEC-1998
15-DEC-1998
15-DEC-1998
                                                                                                                                                         activase) (RA) (Water stress responsive pro
Pinus pinaster (Maritime pine).
Eukaryota; Viridiplantae; Streptophyta; Emb
Spermatophyta; Coniferopsida; Coniferales;
NCBI_TaxID=71647;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Unknown protein from 2D-page of liver tissue
(Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -!- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PROTEIN IS: 6.0, ITS MW IS: 12.5 kDa. SWISS-ZDPAGE, P99031; MOUSE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; M58516; AAA16528.1; -.
EMBL; X02872; CAA26630.1; -.
EMBL; X60827; CAA43217.1; -.
EMBL; X65462; CAA46454.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOUSE
"Water-deficit-responsive proteins Plant Mol. Biol. 38:587-596(1998). [2]
                                                            MEDLINE=98418576; PubMed=9747804;
Costa P., Bahrman N., Frigerio J.-M.,
                                                                                                                                                                                                                                                                                                                                                        P81084;
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Mammalia; Eutheria;
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Leader peptide; Antibiotic resistance; Plasmi

SECUENCE 9 AA; 1074 MW; 5D9CAB5AAB05B333
                                                                                                     TISSUE=Needle;
                                                                                                                           SEQUENCE
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Rodentia;
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Binz P.-A., I
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Pred. No.
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                                         maritime
                                                            Kremer A.,
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                                                                                                                                                                                                                                                                       (RuBisCO
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GenCore version 5.1.3 Compugen Ltd.

OM protein - protein search, using sw model

Run on: February 8, 2003, 11:01:52; Search time 28 Seconds (without alignments) 73.588 Million cell updates/sec

Title: Perfect score: Sequence: US-09-685-010-31 46 1 KLKSQLVKRK 10

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 671580 seqs, 206047115 residues

Total number of hits satisfying chosen parameters: 1224

Minimum DB seq length: 0 Maximum DB seq length: 10

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

SPTREMBL 21:*

SPTREMBL 21:*

Sp archea:*

Sp bacteria

Sp fungi:*

Sp human:*

Sp invert

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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ALIGNMENTS

RESULT 078226 ID 0 AC 0 DT 0	ве ма Оу Db	STREET REPRESENTATION OF STREET	RESULT Q84273 ID Q AC Q DT 0 DT 0 DT 0 DT 0
ULT 2 .226 .078226 PRELIMINARY; PRT; 9 AA. .078226; .01-NOV-1998 (TrEMBLrel. 08, Created)	Similarity 75.0%; Pred. No. 6.7e+05; 3; Conservative 1; Mismatches 0; K 10 1	Human papillomavirus type 25. Viruses; dsDNA viruses, no RNA stage; Papillomaviridae; Papillomavirus. NCBI_TaxID=10609; [1] SEQUENCE FROM N.A. MEDILINE=88089511; PubMed=2826651; Krubke J., Kraus J., Delius H., Chow L., Broker T.R., Iftner T., Pfister H.; "Genetic relationship among human papillomaviruses associated with benign and malignant tumours of patients with epidermodysplasia verruciformis:"; J. Gen. Virol. 68:3091-3103(1987). EMBL; D00205; BAA00144.1; SEQUENCE 8 AA; 999 MW; 7F43240324033058 CRC64; Sequery March 39.1%. Score 18. DR 12: Length 8:	1 84273 PRELIMINARY; 84273; PRELIMINARY; 84273; (TrEMBLrel. 0: 1-NOV-1996 (TrEMBLrel. 0: 1-NOV-1998 (TrEMBLrel. 0: 1-NOV-1998 (TrEMBLrel. 0:

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Q15897;
01-NOV-1996
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Q94II9;
01-DEC-2001
ol-NOV-1996 (TrEMBLrel. 01, Created)
ol-NOV-1996 (TrEMBLrel. 19, Last sequence update)
ol-DEC-2001 (TrEMBLrel. 19, Last annotation update)
(clone XP6A11A) (Fragment).
Homo sapiens (Human).
Eukarbota, Metazoa; Chordata; Craniata, Vertebrata;
Mamm. Metazoa; Chrimates; Catarrhini; Hominidae
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Eukaryota; Magnoliophyta; Liliopsida; Poales; Poaceae; PACC clade;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACC clade;
Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "A maize rl gene is regulated splicing of its leader."; Submitted (APR-2001) to the EN
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update
Anthocyanin regulator R-sc protein (Fragment).
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"Nucleotide sequence analysis of chromosomes.";
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MEDLINE=88084418; PubMed=3692165;
Golubic M., Budimir O., Schoepfer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
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EMBL; L38590; AAA57294.1; -.
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Mammalia; Eutheria; Rodentia;
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3; Conserv
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  Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Matches 3; Conserv
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Q9YIRO;
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SEQUENCE 7 AA; 980 MW;
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NON TER
SEQUENCE
Human adenovirus type 7. Viruses; dsDNA viruses,
                                                           PVI core
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Submitted (DEC-2000)
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Crawford-Miksza L.K., Nang R.N., Schnurr I
"Strain variation in adenovirus serotypes
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J. Clin. Microbiol.
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Coolbaugh M.I., Chinault C.A.,
Caskey C.T.H.;
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1 VKRR 4
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Baldini
stage; Adenoviridae; Mastadenovirus.
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dini A., Lindsay
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SEQUENCE
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Crawford-Mikeza L.K., Nang R.N., Schnurr D.P.;
"Molecular surveillance of strain variation in adenoviruses acute respiratory disease, AV 4 and AV 7a.";
Submitted (MAX-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AP065064; AAD03659.1; -.
EMBL; AP065063; AAD03655.1; -.
EMBL; AP065063; AAD03656.1; -.
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Crawford-Miksza L.K., Nang R.N., Schmurr D.D.;
"Molecular surveillance of strain variation in adenoviru
acute respiratory disease, AV 4 and AV 7a.";
Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF065068; AAD03668.1; -.
EMBL; AF065066; AAD03664.1; -.
EMBL; AF065067; AAD03666.1; -.
                                                                                      Viruses; ssRNA negative-strand Influenza A viruses.
                                                                                                                                                                        01-NOV-1996 (TrEMBLrel. 01, 01-NOV-1996 (TrEMBLrel. 01, 01-MAR-2002 (TrEMBLrel. 20, Influenza A/udorn/72 (H3n2)
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MEDLINE=83112211;    PubMed=6296449;
                                                             NCBI_TaxID=11320;
                                                                                                                               Influenzavirus
                           SEQUENCE FROM N.A.
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1; Mismatches
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Proc. Natl. Acad. Sci. U.S.A.
EMBL; U04930, AAD13472.1; -.
EMBL; U04931; AAD13476.1; -.
NON TER 1 1 1
NON TER 8 8 8 8 SEQUENCE 8 AA; 902 MW; 922
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01-MAY-1997 (TrEMBLrel. 03, L:
01-MAY-1999 (TrEMBLrel. 10, L:
Growth hormone 2 (Fragment).
GH-2-A OR GH-2-B.
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J. Virol. 45:434-438(1983).
EMBL; J02170; AAA43468.1; -.
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01-MAY-1999 (TrEMBLrel. 10,
01-MAY-1999 (TrEMBLrel. 10,
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MEDLINE=94173893; PubMed=8127856;

MIDDINE=94173893; PubMed=8127856;

MIDDINE=94173893; PubMed=8127856;

MIDDINE=94173893; PubMed=817856;

MIDDINE=94173893; Pide=941894;

MIDDINE=941894; 
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleost
Actinopterygii; Metazoa; Endeostei; Buteleostei;
Projacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
                                                                                                                                                                 Buchnera aphidicola.
Plasmid pBRc1.
Bacteria; Proteobact
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    MEDLINE=99028904; PubMed=9812361;
                                       SEQUENCE FROM N.A.
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STRAIN=K-12;
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STRAIN=B6196428; PubMed=8648624;
Mahdi A.A., Sharples G.J., Mandal T.N., I
"Hôlliday Junction Resolvases encoded by
Escherichia coli K-12 and Phage 32.";
J. Mol. Biol. 257:561-573(1996).
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Orf33, orf151, o
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Escherichia coli
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                                                                                           Bacteria; Proteobacteria;
Escherichia.
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Aphididae.";
PEMS Microbiol. Lett. 168
EMBL; AJ006874; CAA07290.
                                                                SEQUENCE FROM N.A.
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                                      SEQUENCE
                                                              Arai E., Ikeuchi T., Nakamura Y., "Characterization of the translocation breakpoint on chromosome 22q12.2 in a patient with neurofibromatosis type 2 (NF2)."; Hum. Mol. Genet. 3:937-939(1994).
                                                                                   MEDLINE=95038750; PubMed=7951241;
Arai E., Ikeuchi T., Nakamura Y.;
"Characterization of the transloc
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Mammalia; Eutheria;
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Q1-MAY-2000 (TrEMBLrel. 13, Created)
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O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Acetyl-CoA acetyltransferase 2 (Fragment).
ACAT2.
ACAT2.
Monodelphis domestica (Short-tailed grey opossum).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Metatheria; Didelphimorphia; Didelphidae; Monodelphis.
NCBI TaxID=13616;
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MEDILINE=99282512; PubMed=10353914;

MEDILINE=99282512; PubMed=10353914;

Shintani S., O'hUigin C., Toyosawa S., Michalova V., Klein J.;

"Origin of gene overlap. The case of tcpl and acat2.";

"Cenetice 152:743-754(1999).

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ALIGNMENTS

Hyalauronan (HA) binding peptide #10. 13-AUG-2002 ABG60827; ABG60827 standard; Peptide; 10 AA. (first entry)

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KW B Tissue disorder; response-to-injury process; cell proliferating; hyaluronic acid; HA; receptor for hyalauronan-mediated motility; RHAMM; inflammatory neurological disorder; Parkinson's disease; RHAMM; inflammatory neurological disorder; Parkinson's disease; halzheimer's disease; arthritis; multiple sclerosis; gastritis; nephritis; inflammatory dermatosis; psoriasis; inflammatory bowel disease; stenosis; restenosis; cancer; kidney fibrosis; inflammatory lung disease; stenosis; restenosis; cancer; kidney fibrosis; inflammatory response disease; lupus; cardiovascular disease; atherosclerosis; wound; scar; diabetes; lupus; cardiovascular disease; atherosclerosis; wound; scar; diabetes; lissue transplantation; stroke; inflammatory response; fibrotic response; 05-OCT-2000; 2000WO-IB01534. 11-APR-2002 WO200228415-A1. Synthetic. septic shock; medical implant; Acquired immunodeficiency syndrome; AIDS; hepatitis, myocardial fibrosis; hepatic fibrosis; chronic cystitis; acute mastit thyroiditis; retinopathy

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RESULT 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention describes a method of treating a tissue disorder associated CC with response-to-injury process or proliferating cells in a patient, CC comprising administering polypeptide (I) which binds hyaluronic acid (HA), an antibody which binds one of domains D1-D5 of Receptor for CC hyalauronan-mediated motility (RHAMM), a polypeptide fragment encoding CC any of D1-D5 of RHAMM, or a vector which expresses antisense RHAMM, CC antibodies or a polypeptide fragment. The method is useful for treating a gatient with an inflammatory neurological disorder such as parkinson's CC disease, Alzheimer's disease, arthritis including rheumatoid arthritis, costecarthritis, multiple sclerosis, inflammatory dermatosis (psoriasis), CC inflammatory bowel disease, stenosis or restenosis, cancer, kidney CC fibrosis, inflammatory lung disease (e.g. emphysema, asthma, cystic CC disease (e.g. atherosclerosis), and wound especially surgical excision CC adhesions, to prevent scar and also for treating or preventing disbetes CC mellitus. The method is also useful for treating tissue transplantation (e.g. skin grafts), stroke, inflammatory responses or fibrosic cresponse CC associated with medical implants such as hIDS, myocardial and hepatic CC fibrosis, chronic cystitis, acute mastitis, saptritis, nephritis, core represents a hyslauronan (HA) binding peptide used in the method of treating a tissue disorder described in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                           Hyaluronan binding protein; HA; RHAMM; mediated motility; wound; healing; diagnosis; treatment; cell locomotion; tumour invasion; birth defects; inflammatory disorder; Alzheimer's disease; dementia; Parkinson's diseases; Huntington's disease; AIDS; diabetes; auto; immune diseases; corneal dysplasia; hypertrophy; surgery; burns; strokes; multiple sclerosis; depression; schizophrenia; CNJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Treating tissue disorder associated with response-to-injury process or proliferating cells in mammals, e.g. fibrosis, inflammation, by administering a compound that alters activity of transition molecules within a cell
09-APR-1992;
                                                              13-APR-1993;
                                                                                                                                                                                                    WO9321312-A.
                                                                                                                                                                                                                                                                  contraception;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hyaluronan receptor binding motif (claimed).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR46551;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR46551 standard; Protein; 10
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   92GB-0007949
                                                                 93WO-CA00158
                                                                                                                                                                                                                                                                  in vitro fertilisation; embryo development.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
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Pred. No. 0.026;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hyaluronan receptor; receptor for hyaluronic acid mediated motility; RHAMM; glycosaminoglycan; binding domain; mouse; oncogene; treatment growth factor; cell locomotion disorder; dementia; detection;
Isolated human receptor for hyaluronic acid mediated motility - used to develop products for treating e.g. tumours, inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                            WPI; 1997-512715/47.
                                                                                                                                               Entwistle J,
                                                                                                                                                                                                                                                                                              10-APR-1996;
                                                                                                                                                                                                                                                                                                                                                    10-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                16-OCT-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse RHAMM binding domain 2 consensus motif peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW39164 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA encoding hyaluronan receptor - used to produce proteins and antibodies for alteration of cell locomotion
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                                                                                                                                                                                                                                      (MANI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inflammatory
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                                                                                                                                                                                                                                      MANITOBA CANCER TREATMENT
                                                                                                                                                                                                           AIND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 9; Conserv
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                                                                                                                                                                                                              MANITOBA.
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                                                                                                                                            Turley EA;
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Pred. No. 0.097;
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in vitro fertilisation and
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disorders,

dementia, AIDS, diabetes and auto-immune diseases

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Matches
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                                                                                                                            Isolated human receptor for hyaluronic acid mediated motility to develop products for treating e.g. tumours, inflammatory disorders, dementia, AIDS, diabetes and auto-immune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hyaluronan receptor; receptor for hyaluronic acid mediated motility; RHAMM; glycosaminoglycan; binding domain; rat; oncogene; treatment; growth factor; cell locomotion disorder; dementia; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cell surfaces via specific protein receptors, e.g. RHAMM, that mediate many biological effects. The RHAMM/Hyaluronic acid interaction is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This peptide represents a motif found in a binding domain of mouse hyaluronan receptor corresponding to amino acid position 424-433. Treceptor is also known as the receptor for hyaluronic acid mediated
                                                                   Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rat RHAMM binding domain 2 consensus motif peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW39168 standard; peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                              WPI; 1997-512715/47.
                                                                                                                                                                                                                                                                                                                           Entwistle J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-OCT-1997.
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mes 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                          MANITOBA CANCER TREATMENT & RES FOUND. UNIV MANITOBA.
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                                                               Fig 1; 66pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disorder; autoimmune disease; diagnosis; prognosis.
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                                                                                                                                                                                                                                                                                                                           Turley EA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 43; DB 18; Length 10; Pred. No. 0.097;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and prognosis.
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This peptide represents a motif found in a binding domain of rat

The sequence is that of a binding motif fragment of the hyaluronan receptor (HARC). HARC is down regulated in normal cells and is on expressed in situations where cell motifity is desired, e.g. in wound healing, in response to growth factors and in chemotaxis by white blood cells. HA may be used for diagnosis and treatment of

and is only

diseases involving

locomotion, e.g. tumour invasion,

DNA encoding hyaluronan receptor - antibodies for alteration of cell]

locomotion

used to produce proteins

WPI; 1993-351722/44

Claim 17; Page 57; 88pp; English.

09-APR-1992;

92GB-0007949.

(MANI-) MANITOBA CANCER TREATMENT & RES (UYMA-) UNIV MANITOBA.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                birth defects; inflammatory disorder; Alzheimer's disease; dementia; Parkinson's diseases; Huntington's diseases, AIDS, diabetes; auto; immune diseases; corneal dysplasia; hypertrophy; surgery; burns; strokes; multiple sclerosis; depression; schizophrenia; CNJ; contraception; in vitro fertilisation; embryo development.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hyaluronan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hyaluronan receptor binding motif (claimed).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-APR-1994 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR46550;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-APR-1993;
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8; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              binding protein; HA; RHAMM; mediated motility; wound; iagnosis; treatment; cell locomotion; tumour invasion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10 AA;
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Pred. No. 0.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tissue disorder; response-to-injury process; cell proliferating; hyaluronic acid; HA; receptor for hyalaurona-mediated mottlity; RHAMM; inflammatory neurological disorder; Parkinson's disease; Alzheimer's disease; arrhritts; multiple sclerosis; gastritts; nephritis; inflammatory dermatosis; psoriasis; inflammatory bowel disease; stenosis; restenosis; cancer; kidney fibrosis; inflammatory lung disease; emphysema; asthma; cystic fibrosis; obesity; obesity related disease; lupus; cardiovascular disease; atherosclerosis; wound; scar; diabetes; tissue transplantation; stroke; inflammatory response; fibrotic response; medical implant; Acquired immunodeficiency syndrome; AIDS; hepatitis; myocardial fibrosis; hepatic fibrosis; chronic cystitis; acute mastitis; septic shock; thyroiditis; retinopathy.
      with response-to-injury process or proliferating a tissue disorder associat with response-to-injury process or proliferating cells in a patient, comprising administering a polypeptide (I) which binds haluronic acid (HA), an antibody which binds one of domains D1-D5 of Receptor for hyalauronan-mediated motility (RHAMM), a polypeptide fragment encoding apy of D1-D5 of RHAMM, or a vector which expresses antisense RHAMM, and the processes antisense RHAMM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     defects, acute and chronic inflammatory disorders, Alzheimer; s and other forms of dementia, AIDS, diabetes, autoimmune diseases, corneal dysplasias and hypertrophies, burns, surgical incisions and adhesions, strokes, multiple sclerosis, depression/schizophrenia related to neuronal growth and pain states involving nerve sprouting; also in CNJ and spinal cord regeneration, contraception, in vitro fertilisation and
                                                                                                                               Disclosure; Page
                                                                                                                                                                             proliferating administering
                                                                                                                                                                                                             Treating
                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                              Turley EA, Cruz TF
                                                                                                                                                                                                                                                                                                                                               05-OCT-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cellular response to injury associated peptide
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so AAR46548-51 and AAR43563.
                                                                                                                                                                                                                                                                                                             TRANSITION THERAPEUTICS & DIAGNOSTICS
                                                                                                                                                                           tissue disorder associated with response-to-injury process ating cells in mammals, e.g. fibrosis, inflammation, by ering a compound that alters activity of transition molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
 or a
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polypeptide
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88.9%;
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Pred. No.
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 The method is useful
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treating
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 osteoarthritis, multiple scierosis, inflammatory dermatosis (psoriasis), inflammatory bowel disease, stenosis or restenosis, cancer, kidney fibrosis, inflammatory lung disease (e.g. emphysema, asthma, cystic fibrosis), obesity or obesity related diseases, lupus, cardiovascular disease (e.g. atherosclerosis), and wound especially surgical excision adhesions, to prevent scar and also for treating or preventing diabetes mellitus. The method is also useful for treating tissue transplantation (e.g. skin grafts), stroke, inflammatory responses or fibrotic response associated with medical implants such as hip implants, vascular wraps an (e.g. chronic cystitis, acute mastitis, gustritis, mephritis, acute mastitis, gastritis, sephritis, hepatitis, septic shock, thyroiditis, and retinopathy. This sequence represents a peptide associated with the method of treating tissue
                                                                                                     Protein conjugates that selectively target certain tissues and organs useful for treating and preventing various diseases, comprises glucose-aminoglycan-targeting domain conjugated to a therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hyaluronic acid-binding peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAU11442;
                                                                                                                                                                                                                                          20-APR-2000; 2000US-198613P
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                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                  conjugate; rheumatoid arthritis; scleroderma; liver fibrosis; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hyaluronic acid binding protein; RHAMM; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAU11442 standard; Peptide; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          patient with an inflammatory neurological disorder such as Parkinson's
                                                                                                                                                                                                                                                                                                                         WO200180899-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KLRSQLVKR
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8; Conserv
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                                                                                                                                                                                                                  CANGENE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                 for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alzheimer's disease, arthritis including rheumatoid nritis, multiple sclerosis, inflammatory dermatosis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                HA mediated mobility;
                                                                  109;
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Pred. No.
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7.8e+05;
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The invention relates to a conjugate comprising an hyaluronic acid binding protein e.g. RHAMM (receptor of HA mediated mobility) or promise with, or coupled to a polypeptide conjugated to a therag agent, and the polynucleotides encoding them. Also included is a me

y) or peptide therapeutic is a method

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                                                                                                                                                     Isolated human receptor for hyaluronic acid mediated motility to develop products for treating e.g. tumours, inflammatory disorders, dementia, AIDS, diabetes and auto-immune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hyaluronan receptor; receptor for hyaluronic acid mediated motility; RHAMM; glycosaminoglycan; binding domain; human; oncogene; treatment; growth factor; cell locomotion disorder; dementia; detection;
                                                                               Claim 7; Fig 1; 66pp; English.
                                                                                                                                                                                                                                                                                                                                          WPI; 1997-512715/47.
                                                                                                                                                                                                                                                                                                                                                                                                                         Entwistle J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-APR-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MANI-) MANITOBA CANCER TREATMENT & RES FOUND. (UYMA-) UNIV MANITOBA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inflammatory disorder; autoimmune disease; diagnosis; prognosis.
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This peptide represents a motif found in a binding domain of human

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Best Local Similarity
""" hes 6; Conserv?
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                                                                                                                                                                                                                                                                                                                                   17-JUL-1986;
02-SEP-1986;
20-NOV-1986;
27-JUL-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Region
Recombinant bovine rhinotracheitis virus expressing antigenic polypeptide - useful as multivalent vaccine, esp. against parainfluenza or rotavirus
                                                                                                           N-PSDB;
                                                                                                              WPI; 1996-340868/34.
N-PSDB; AAT31911.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Swine parvovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-SEP-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR99730;
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                                                                                                                                                                                             Chiang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-JUL-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US5506128-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vector; gpX secretion signal; S-PRV-040.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pseudorabies virus; PRV; swine parvovirus; herpesvirus; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Plasmid 167-72.54G Junction C-encoded peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-SEP-1985;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-SEP-1985;
                                                                                                                                                                                                                                                                                                           17-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-JAN-1986;
                                                                                                                                                                                                                                                       (PRUT-)
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                                                                                                                                                                                                                                                    PRUTECH
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86US-0933107.
87US-0078519.
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85US-0773430.
86US-0823102.
86US-0887140.
                                                                                                                                                                                                                                                       RES
                                                                                                                                                                                                                                                                                                           93US-0078873.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                85US-0773430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "capsid protein C-terminal region'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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60.0%;
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Pred. No.
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Matches
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                                                                                                                                                                                                                                                                          02-SEP-1986;
20-NOV-1986;
27-JUL-1987;
17-JUN-1993;
Plasmid 244-25.3D contains an E. coli beta-galactosidase-swine parvovirus A capsid fusion gene as an Xbal Eragment in vector pSPr The sequences of the junctions between 3 segments of the fusion gene and the vector are given in AAT13801, AAT13802, AAT31902 and AAT31903. The peptide (AAR99725) encoded by Junction D (AAT31903) comprises the C-terminal region of capsid A. The fusion gene was inserted into the repeat region of PRV, giving recombinant virus
                                                                                                                              Recombinant bovine rhinotracheitis virus expressing antigenic polypeptide - useful as multivalent vaccine, esp. against parainfluenza or rotavirus
                                                                                                       Example
                                                                                                                                                                                                                        Chiang
                                                                                                                                                                                                                                                                                                                                          06-SEP-1985;
27-JAN-1986;
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                                                                                                                                                                                                                                                                                                                                                                                             06-SEP-1985;
                                                                                                                                                                                                                                                                                                                                                                                                                      09-APR-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Swine parvovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pseudorabies virus; PRV; swine parvovirus; herpesvirus; vaccine; capsid; beta-galactosidase; vector; S-PRV-065.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Plasmid 244-25.3D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR99725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR99725 standard; Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Plasmid 167-72.54G contains a pseudorables virus (PRV) gpX secretion signal-swine parvovirus A capsid fusion gene as an XbaI fragment in vector pSP65. The fusion gene was prepd. in 2 segments. The junction (AAT31911) between the segment 2 and the vector includes a region coding for the C-terminal region (AAR89729) of the capsid protein. The fusion gene was inserted into the repeat region of PRV, glving recombinant virus S-PRV-040, which can be used as a vaccine to protect swine against parvovirus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 28;
                                                                                                                                                                                                                                                  (PRUT-)
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                                                                                                                                                                                   1996-340868/34.
DB; AAT31903.
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5; Conserv
                                                                                                     26; Fig 40B; 114pp; English.
                                                                                                                                                                                                                                                   PRUTECH RES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                          88US-0225032.

85US-073430.

86US-0823102.

86US-0887140.

86US-0902887.

86US-0932107.

87US-0078519.

93US-0078873.
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                                                                                                                                                                                                                                                 & DEV PARTNERSHIP.
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71.4%;
                                                                                                                                                                                                                          MacDonald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      D-encoded
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Pred. No. 7.8e
1; Mismatches
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7.8e+05;
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RESULT 12
ABG35074
ID ABG35
XX
AC ABG35
XX

ABG35074 standard;

Peptide;

9

ABG35074;

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RESULT 11
AAG95218
ID 95218
ID 95218
IX AAG95
XX AAG95
XX IN Human
XX Human
XX Homo
XX Homo
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                                                                                                 Matches
                                                                                                                                          Query Match
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Best Local
                                                                                                                                                                                                                                         The invention relates to a set of complementary peptide ligands generated from the human genome. The complementary peptides interact with their relevant target proteins encoded in the human genome. They can be used as reagents in drug discovery and as lead ligands to facilitate drug design and development. The present sequence is a complementary peptide provided in the specification.
                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                              A set of peptide ligands consisting of specific complementary peptides to proteins encoded by genes of the human genome, useful in an assay for screening and identifying of one or more novel peptides which are drug candidates or pro-drugs -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         parvovirus
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                                                                                                                                                                                                                                                                                                                                                                                                                 Example 4; Page 243; 646pp; English.
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Pred. No. 3.7e
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Pred. No. 7.8e
1; Mismatches
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                                                                                                                        3.7e+02;
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7.8e+05;
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                                                                                                                                          Length 10;
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                                                                                                                                                                                                                                                                                                                                                                     This invention relates to a novel method for identifying disease targeting peptides. The method comprises exposing a sample from an organ, tissue or cell type of interest, to a phage display library and recovering phage bound to the sample (the phage expresses targeting peptides). The peptides identified by the method of the invention may have cyrostatic, immunosuppressive, anti-inflammatory, antiarthritic, antidiabetic, antibacterial and antiviral activities. The methods and composition are useful for identifying targeting peptides and one or more receptors for a targeting peptide. The targeting peptides are used for selective delivery of therapeutic agents, including gene therapy vectors and fusion proteins, to specific organs, tissues, or cell types in subject. The targeting peptide may also be used for treating diseases such as diabetes mellitus, inflammatory diseases, arthritis, atheroselerosis, cancer, autoimmune disease, bacterial and viral infections and Hodgkin's disease. The present sequence represents a targeting peptide of the invention.
                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                            Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying targeting peptides useful for treating e.g. diabetes mellitus, inflammatory diseases, cancer, or autoimmune diseases, comparises exposing a sample to a phage display library and recovering phage bound to the sample -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-SEP-2000; 2000US-231266P.
17-JAN-2001; 2001US-0765101.
                                                                                                 AAR90268
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                                  Ion-channel
                                                                10-ЛП-1996
                                                                                                                             AAR90268 standard; peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200220722-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Targeting peptide; cancer; Hodgkin's disease;
                                                                                                                                                                                                                                                                            Local Similarity
108 5; Conserv
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                                                                                                                                                                                                                                            6 LVKRK 10
                                                                                                                                                                                                               LVKRK 7
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                              forming
forming peptide, lipophilic; N-terminal modification,
                                                                                                                                                                                                                                                                              Conservative
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100.0%; Pr
                              peptide #129 with lipophilic N-terminal group.
                                                              entry)
                                                                                                                                                                                                                                                                                             Score 23;
Pred. No.
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hes 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               These ionophores are known to have a broad range of potent antibiotic activity against microorganisms including gram-positive and gram-negative bacteria, fungi, viruses, protozoa and parasites. N-terminal modification (pref. mono-substr. by octanoy1) to produce an ion-channel forming peptide having a lipophilic N-terminus increases the biological activity of the peptides against target cells, viruses and virally-infected cells, compared to peptides substd. with an acetyl group at the N-terminus. Compositions comprising the peptides with lipophilic modifications are claimed for inhibiting growth of a target cell, virus or virally-infected cell.
                Sepsis; septic shock; Pseudomonas aeruginosa; cystic fibrosis; antimicorobial; antiviral; antibacterial; antifungal; antitumour; antiparasitic; spermicide; preservative; sterilant; disinfectant; wound healing; burn; skin infection; eye infection; solid tumour; leukaemia; non-small cell lung cancer; adenocarcinoma; plant infection periodontal disease; plaque; gingivitis; caries; Streptococcus mut
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  magainin; inhibition; cell growth; viral replication; ionophore; membrane permeability; antimicrobial; anti-bacterial; antibiotic; anti-fungal; anti-viral; spermicidal; anti-tumour; anti-parasitic.
                                                                                                                                                                                Peptide used to make biologically active peptides.
                                                                                                                                                                                                                                11-MAY-1999
                                                                                                                                                                                                                                                                                                                      AAY10768 standard; peptide; 10 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 30;
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lipophilic gps. - useful e.g. as antiviral, antibacterial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kari UP,
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octanoyl"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 16; Le. 5.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 10;
                       Streptococcus mutans.
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                                             plant infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 15
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                                                                                                                                                                                                                                      Biologically active peptide; toxicity; antimicrobial; anti-tumour; methane sulphonate derivative; wound healing; burn; therapy; sepsis; eye infection; cyst; spore; trophozoite; tumour; lung infection; cystic fibrosis; septic shock; bacterial endotoxin; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Treating sepsis or septic shock with N-modified ion-channel forming peptide - or its methane sulphonate derivative of reduced toxicity, also generally useful as antimicrobial and antitumour agents
Modified-site
                                                              Modified-site
                                                                                                                                                     Unidentified
                                                                                                                                                                                                                                                                                                                                                                                              Biologically
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                                                                                                                                                                                                             immunosuppressive.
   /note= "Linked to octanoyl group"
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                                                                                       Location/Qualifiers
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3; Mismatches
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5.7e+02;
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Search completed: February
Job time : 36 secs
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                                                                                                                                                                                                      cc toxicity and methods of preparing them. The peptides and proteins of the invention have improved antimicrobial and anti-tumour activity while exhibiting reduced toxicity. The method of reducing toxicity convolves the formation of related methane sulphonate derivatives or analogues. The method is useful for reducing the toxicity of an cumsubstituted peptide or an N-terminal substituted peptide which is cutilised in promoting or stimulating healing of a wound in a host, cutilised in promoting or stimulating healing of a wound in a host, caused by bacteria or fungi, in killing cysts, spores, or trophozoites caused by bacteria or fungi, in killing cysts, spores, or trophozoites confinection-causing organisms, and may also be employed in the creatment of tumours, serious lung infections such as those occurring in cystic fibrosis, for treating sepsis, septic shock, and other related ailments, and for neutralising bacterial endotoxins. The present sequence is a biologically active peptide of the invention.
                                                                                                                                    Matches
                                                                                                                                                  Query Match
Best Local
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01-JUN-1992;
05-JUN-1995;
15-JUL-1997;
                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Reducing toxicity of unsubstituted or N-terminal substituted peptide having antimicrobial and antitumor activity useful in treating e.g. infections and tumor, by forming methane sulfonate derivative or analog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-350076/38.
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                                                                                                                                                Local Similarity
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95US-0465330.
97US-0893006.
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                 2003, 11:03:09
                                                                                                                                    3; Mismatches
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Minimum
Maximum
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Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                              Published
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/cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
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/cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
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(c) 1993 - 2003 Compugen Ltd
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

19	18	17	16	15	14	13	12	11	10	9	89	7	6	v	4	ω	N	<u>, 1</u>	Result No.	
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US-09-878-603-21	US-09-752-533-2	US-09-798-116-19	US-09-789-996-42	US-09-867-852-207	US-09-867-852-205	US-10-100-057-15	US-09-520-698-6	US-09-520-698-5	US-09-520-698-1	US-09-805-301-1	US-09-878-603-20	US-09-243-079-26	US-10-147-454-6	US-09-877-804-22	US-09-819-308-46	US-09-779-308-481	US-09-779-308-182	US-09-878-603-14	ID	
Sequence 21, Appl	Sequence 2, Appli	Sequence 19, Appl	Sequence 42, Appl	Sequence 207, App	Sequence 205, App	Sequence 15, Appl	٠ م	Sequence 5, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 20, Appl	Sequence 26, Appl	Sequence 6, Appli	Sequence 22, Appl	Sequence 46, Appl	Sequence 481, App	Sequence 182, App	Sequence 14, Appl	Description	

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20
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US-10-211-207-15	US-10-211-207-1	US-09-178-286-7	US-09-957-909-2	US-09-883-825-2	US-09-243-079-48	US-09-243-079-27	US-09-989-789-222	US-09-989-789-216	US-09-989-789-214	US-09-989-789-213	US-09-989-789-211	US-09-989-789-208	US-09-779-308-63	US-09-752-533-7	US-09-752-533-4	US-09-124-280A-16	US-09-838-785-24	US-09-779-308-211	US-09-905-083-47	US-09-918-243-47	US-09-752-533-3	US-09-834-765-331	US-09-834-765-219	US-09-834-765-18	US-09-765-527-220
Sequence 15, Appl	بر	Sequence 7, Appli	.~			٠.		o .	o.	٠.	o	e 208,	_	7,	Sequence 4, Appli	Sequence 16, Appl	e 24, 1	211,	e 47	e 47	w	33	21	~	Sequence 220, App

ALIGNMENTS

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; TYPE: PRT; ORGANISM: Homo sapiens US-09-878-603-14
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US-09-779-308-182
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                 Sequence 182, Application US/09779308
Patent No. US20020150972A1
GENERAL INFORMATION:
APPLICANT: Mary Faris
APPLICANT: Daniel E.H. Afar
APPLICANT: Pia M. Challita-Eid
APPLICANT: Rene S. Hubert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 14, Application US/09878603
Patent No. US20020165138A1
GENERAL INFORMATION:
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Best Local Similarity 44.4%;
Matches 4; Conservative
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SEQ ID NO 14
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Ward, Peter A.
APPLICANT: Huber-Lang, Markus
APPLICANT: Huber-Lang, Markus
APPLICANT: Sarma, Vidya
APPLICANT: Czermak, Boris
TITLE OF INVENTION: Compositions and Methods for the Treatment of Sepsis
FILE REFERENCE: UM-03783
CURRENT APPLICATION NUMBER: US/09/878,603
CURRENT FILING DATE: 2001-66-11
PRIOR APPLICATION NUMBER: 09/387,671
PRIOR FILING DATE: 1999-08-31
PRIOR FILING DATE: 1999-08-31
PRIOR FILING DATE: 1999-08-31
PRIOR FILING DATE: 1999-08-31
PRIOR SECOND NOS: 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                      Elana Levin
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Pred. No. 1.1e+05;
2; Mismatches 3
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APPLICANT: SLEAR LEVIN
APPLICANT: Steve Chappell Mitchell
APPLICANT: Steve Chappell Mitchell
APPLICANT: Steve Chappell Mitchell
APPLICANT: ANG Jakobovits
TITLE OF INVENTION: 349307: A TISSUE SPECIFIC PROTEIN
TITLE OF INVENTION: HIGHLY EXPRESSED IN PROSTATE CANCER
FILE REFERENCE: 129 4 USUI
CURRENT APPLICATION NUMBER: US/09/779,308
CURRENT FILING DATE: 2001-02-08
PRIOR APPLICATION NUMBER: 60/181,020
PRIOR APPLICATION NUMBER: 60/181,020
PRIOR FILING DATE: 2000-02-08
NUMBER OF SEQ ID NOS: 718
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 481
LENGTH: 9
TYPE: PRT
ORGANISM: Homo Sapiens
US-09-779-308-481
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                                                                              GENERAL INFORMATION:
APPLICANT: No. US20020019040A1eborn, Mathieu
APPLICANT: Danen-van Oorschot, Astrid
APPLICANT: Rohn, Jennifer
                                                                                                                                                                  Sequence 46, Application US/09819308 Patent No. US20020019040A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Mary Faris
APPLICANT: Daniel E.H.
APPLICANT: Dia M. Chal
APPLICANT: Rene S. Hub
APPLICANT: Elana Levin
APPLICANT: Steve Chapp
APPLICANT: Aya Jakobov
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CURRENT APPLICATION NUMBER: US/09/779,308
CURRENT FILING DATE: 2001-02-08
PRIOR APPLICATION NUMBER: 60/181,020
PRIOR FILING DATE: 2000-02-08
NUMBER OF SEQ ID NOS: 718
SOPTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 182
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Best Local :
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CURRENT APPLICATION NUMBER: US/09/819,308
                                     TITLE OF INVENTION: APOPTIN-ASSOCIATING PROTEIN FILE REFERENCE: 2906-4820US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: 34P3D7: A TISSUE SPECIFIC PROTEIN TITLE OF INVENTION: HIGHLY EXPRESSED IN PROSTATE CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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ORGANISM: Homo Sapiens
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Local Similarity 44.4%;
hes 4; Conservative
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Pia M. Challita-Eid
Rene S. Hubert
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Aya Jakobovits
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                                                                                                                                                                                                                                                                                                                                                                                                 Score 20; DB 10; Length 9; Pred. No. 1.1e+05;
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Pred. No.
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APPLICANT: MCFAILAND, Neith C.
APPLICANT: Segalo, Deborah L.
APPLICANT: Segalo, Deborah L.
APPLICANT: Seeburg, Peter H.
TITLE OF INVENTION: Glycoprotein Hormone Receptor Molecules
FILE REFERENCE: PO576F1C2
CURRENT PILING DATE: 2001-09-04
PRIOR APPLICATION NUMBER: US/09/877,804
CURRENT FILING DATE: 1994-03-07
PRIOR APPLICATION NUMBER: US 08/207,814
PRIOR APPLICATION NUMBER: US 07/781,153
PRIOR FILING DATE: 1994-03-07
PRIOR APPLICATION NUMBER: US 07/781,683
PRIOR APPLICATION NUMBER: US 07/347,683
PRIOR FILING DATE: 1989-05-05
NUMBER OF SEQ ID NOS: 22
SEQ ID NO 22
                APPLICANT: COLOMBO, Irma
APPLICANT: RONCHI, Severino
APPLICANT: BARTORELLI, Alberto
APPLICANT: BUMELIS, Viadas Algirdus
TITLE OF INVENTION: Recombinant Protein and It
FILE REFERENCE: 0471-0273P
CURRENT APPLICATION NUMBER: US/10/147,454
CURRENT FILING DATE: 2002-05-17
NUMBER OF SEQ ID NOS: 9
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TYPE: PRT
; ORGANISM: Sequence homology analysis
US-09-819-308-46
                                                                                                                                                                                                                                                                                                                                  US-10-147-454-6
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US-09-877-804-22
                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: NAKTINIS, Vytautas
APPLICANT: CONCAS BENEVELLI, Daniela
APPLICANT: BERRA, Bruno
APPLICANT: COLOMBO, Irma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 43.5%;
Best Local Similarity 37.5%;
Matches 3; Conservative
                                                                                                                                                                                                                                                                                      Sequence 6, Application US/10147454 Publication No. US20020193308A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PA
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Nikolics, Karoly APPLICANT: McFarland, Keitl
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NUMBER OF SEQ ID NOS: 46
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                             1 RAELYRRK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 KSQLVKRK 10
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Pred. No. 2.9e+02;
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Pred. No. 2.
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                                                                                                   Its Use
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                                                                                                 Therapy and Diagnostics
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US-09-243-079-26
; Sequence 26, Application US/09243079
; Patent No. US20020081566A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 29928-PCT-USA-I
CURRENT APPLICATION NUMBER: US/09/243,079
CURRENT FILING DATE: 1999-02-02
PRIOR APPLICATION NUMBER: 08/335,733
PRIOR FILING DATE: 1994-11-10
PRIOR APPLICATION NUMBER: PCT/IT93/00049
PRIOR FILING DATE: 1993-05-10
PRIOR APPLICATION NUMBER: RM92A/000350
PRIOR APPLICATION NUMBER: RM92A/000350
PRIOR FILING DATE: 1992-05-11
NUMBER OF SEQ ID NOS: 89
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Best Local S
Matches 4
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SEQ ID NO 26
LENGTH: 8
TYPE: PRT
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                                    SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 20
LENGTH: 9
                                                                                                                                                                                                                                                                                                           Patent No. US20020165138A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                 Sequence 20,
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                                                                                      APPLICANT: Czermak, Boris
TITLE OF INVENTION: Compositions and Methods for the Treatment of Sepsis
FILE REFERENCE: UM-03783
CURRENT APPLICATION NUMBER: US/09/878,603
CURRENT FILING DATE: 2001-06-11
PRIOR APPLICATION NUMBER: 09/387,671
PRIOR FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 74
                                                                                                                                                                                                                       APPLICANT: Huber-Lang, Markus
APPLICANT: Sarma, Vidya
APPLICANT: Czermat
                                                                                                                                                                                                                                                                   APPLICANT: Ward, Peter A. APPLICANT: Huber-Lang, M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -09-243-079-26
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TITLE OF INVENTION: HOWOLOGOUS TO HLA
TITLE OF INVENTION: HOWOLOGOUS TO HLA
FILE REFERENCE: 29928-PCT-USA-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Local Similarity 66.7%;
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Pred. No. 1.1e+05;
2; Mismatches 2
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Pred. No. 1.1e+05;
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                                                                                                                              US-09-805-301-1
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Patent No. US20020173456A1
GENERAL INFORMATION:
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                                                                              Query Match
Best Local Similarity
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                                                                                                                                                                                                                          TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
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                              1 KLKSQLVK 8
                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/584,043
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 6.4
SOFTWARE: Word Perfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/805,301
FILING DATE: 12-Mar-2001
CLASSIFICATION: Unknown>
                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 217/189
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: LIPOPHILIC PEPTIDES FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Smith, Louis C.
KLRKRLLR 9
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STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                             TELEPHONE:
                                                                Conservative
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                                                                              41.3%;
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                                                             Score 19; DB 9;
Pred. No. 1.1e+05;
4; Mismatches 1
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Pred. No. 1.1e+05;
2; Mismatches 3
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US-09-520-698-6
Sequence Application US/09520698
Patent No. US20020128175A1
GENERAL INFORMATION:
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                                                                                                                        RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/09520698
Patent No. US20020128175A1
GENERAL INFORMATION:
APPLICANT: Gattadahalli M. Anantharamaiah
TITLE OF INVENTION: Synthetic Peptides that Enhance LDL Uptake
FILE REFERENCE: D6084
CURRENT FILING DATE: 1999-03-17
PRIOR PILING DATE: 1999-03-17
PRIOR FILING DATE: 1998-03-17
RUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                              Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local !
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APPLICANT: Gattadahalli M. Anantharamaiah
TITLE OF INVENTION: Synthetic Peptides that Enhance LDL Uptake
FILE REFERENCE: D6084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Gattadahalli M. Anantharamaiah
TITLE OF INVENTION: Synthetic Peptides that Enhance LDL Uptake
FILE REFERENCE: D6084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/520,698
CURRENT FILING DATE: 1999-03-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 09/271,066
PRIOR FILING DATE: 1998-03-17
                                                                                                                                                                                                                                                                                                                        NAME/KEY: DOMAIN
LOCATION: 134..143
OTHER INFORMATION: rabbit apolipoprotein E receptor binding domain
                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Lepus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: DOMAIN
LOCATION: 141..150
OTHER INFORMATION: minimal arginine-rich human apolipoprotein
OTHER INFORMATION: receptor binding domain
                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH: 10
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les 3; Conserv
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nes 3; Conserv
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Pred. No. 4.4e+02;
4; Mismatches 1
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Pred. No.
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US-10-100-057-15
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US-10-100-057-15
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Best Local Similarity
Matches 3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/520,698
CURRENT FILING DATE: 1999-03-17
PRIOR APPLICATION NUMBER: 09/271,066
PRIOR FILING DATE: 1998-03-17
NUMBER OF SEQ ID NOS: 10
                                                                                                                                               TELEPHONE: (202) 628-5197
TELEPAX: (202) 737-5528
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
NAME/KEY: DOMAIN
LOCATION: 141..150
OTHER INFORMATION: monkey apolipoprotein E receptor binding domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Monkey sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
                                                                                                                                                                                                                            APPLICATION NUMBER: US/10/100,057
FILING DATE: 19-Mar-2002
PRIOR APPLICATION NUMBER: US/09/030,061
APPLICATION NUMBER: US 55,468/1997
FILING DATE: 25-FEB-1998
APPLICATION.NUMBER: UP 55,468/1997
FILING DATE: 25-FEB-1997
ATTORNEY/AGENT INFORMATION:
NAME: BROWDY, ROGER LS.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: GILLISPIE-1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 KLRKRLLR 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SECULIARY AND NEIMARK ADDRESSEE: BROWDY AND NEIMARK ADDRESSEE: A19 Seventh Street, N.W., Suite 300
                 MOLECULE TYPE: peptide FRAGMENT TYPE: N-terminal fragment SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      KURIMOTO, MASASHI
TITLE OF INVENTION: OSTEOCLASTGENIC INHIBITORY AGENT
NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: GILLISPIE, Matthew Todd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KLKSQLVK 8
                                                                                   TOPOLOGY: linear
                                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 20004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: D.C
                                                                                                                             LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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UDAGAWA, No. US20020150555Aluyuki
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Pred. No. 4.4e+02;
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Query Match

41.3%;

Score 19;

DB 12;

Length 10;

66.7%;

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US-09-867-852-205
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APPLICANT: Ausubel, Frederick M.
APPLICANT: Staskawicz, Brian J.
APPLICANT: Brent, Andrew F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 208
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 207, Application US/09867852 Patent No. US20020147324A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 39.1%; Score 18; DB Best Local Similarity 100.0%; Pred. No. 1.7 Matches 4; Conservative 0; Mismatches
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APPLICANT:
APPLICANT:
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CURRENT APPLICATION NUMBER: US/09/867,852

CURRENT FILING DATE: 2001-05-29

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/301,085

PRIOR FILING DATE: EARLIER FILING DATE: 1999-04-28

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/310,912

PRIOR PILING DATE: EARLIER FILING DATE: 1994-09-22

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/227,360

PRIOR FILING DATE: EARLIER FILING DATE: 1994-04-13
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PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/301,085

PRIOR FILING DATE: EARLIER FILING DATE: 1999-04-28

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/310,912

PRIOR FILING DATE: EARLIER FILING DATE: 1994-09-22

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/227,360

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/227,360

PRIOR APPLICATE: BARLIER FILING DATE: 1994-04-13
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APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE FAMILY, PRIMERS, PROBES,
TITLE OF INVENTION: DETECTION METHODS
FILE REFERENCE: 00786/254002
CURRENT APPLICATION NUMBER: US/09/867,852
CURRENT FILING DATE: 2001-05-29
                                                                                                                                                                                          APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE FAMILY, PR
TITLE OF INVENTION: DETECTION METHODS
FILE REFERENCE: 00786/254002
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TYPE: PRT
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2 KLKS 5
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Mindrinos, Michael N.
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Mindrinos, Michael N.
                                                                                                                                                                                                                                                                                                                                                    Katagiri, Fumiaki
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2; Mismatches 0;
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hes 0;
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; NUMBER OF SEQ ID NOS: 208
; SOFTWARE: FRASTSEQ for Windows Version 4.0
; SEQ ID NO 207
; LENGTH: 6
; TYPE: PRT
; ORGANIZM: Arabidopsis thaliana
US-09-867-852-207
밁
                   Ś
                                          Matches
                                                    Query Match
Best Local S
                                                 Local Similarity
                   1 KLKS 4
                                        4;
                                           Conservative
                                                   39.1%;
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                                                    Score 18;
Pred. No.
                                           Mismatches
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1.1e+05;
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Search completed: February Job time : 12 secs

8, 2003, 11:07:43

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Minimum
Maximum
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Maximum Match
Listing first
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Perfect score:
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                               Pending | 1: /cgn2 | 2: /cgn2 | 3: /cgn2 | 4: /cgn2 | 5: /cgn | 6: /cgn | 7: /cgn | 8: /cgn | 9: /cgn | 10: /cg | 11: /cgn | 11: /cgn | 11: /cgn | 12: /cgn | 12: /cgn | 13: /cgn | 14: /cgn | 15: /cgn | 15: /cgn | 16: /cgn | 17: /cgn | 16: /cgn | 17: /cgn | 18: /cgn | 19: /cg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              seq length: 0 seq length: 10
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46
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Pending Patents AA Main:*

(cgn2_6/ptodata/1/paa/USO6_COMB.pep:*
(cgn2_6/ptodata/1/paa/USO6_COMB.pep:*
(cgn2_6/ptodata/1/paa/USO6_COMB.pep:*
(cgn2_6/ptodata/1/paa/USO8_COMB.pep:*
(cgn2_6/ptodata/1/paa/USO81_COMB.pep:*
(cgn2_6/ptodata/1/paa/USO81_COMB.pep:*
(cgn2_6/ptodata/1/paa/USO84_COMB.pep:*
(cgn2_6/ptodata/1/paa/USO85_COMB.pep:*
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Copyright (c) 1993 - 2003 Compugen Ltd.
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1 100%
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

93.5	4 43 93.5 10 5 43 93.5 10	100.0	2 46 100.0 1	1 46 100.0	Result Query No. Score Match Length DB ID
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18 US-09-424-292-6	US-08-318-892B-4 S US-09-169-077-2	3 US-09-978-309A-31) US-09-685-010-31	7 US-08-318-892-4	DB ID
Sequence 6, Appl:	Sequence 4, Appli Sequence 2, Appli	Sequence 31, App	Sequence 31, App.	Sequence 4, Appli	Description

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412-8	US-09-412-863-3187	-10-177-725-	2-404B-	US98-10896-5	-10-006-760	08-474-	08-465-330-	US-08-404-283-129	266-2	765-101-	US-09-424-292-17	PCT-US01-27702A-161	US-09-572-404B-1412	663-	-09-424-292-2	08-318-892E	å	077	292	09-424-292-2	-09-424-292-3	08-318-	92-12	-292-	892E	18-892-11	1-292-3	292-3	-09-424-292-2	08-318-892E	18-89	-09-978-309A-	-09-685-010	18-892B-3	-09-169-077-8	-09-424-292-2	US-09-424-292-15
Sequence 9609, Ap	3187,	e 134,	e 198	50,	e 26	6, A	129	129, 1	e 202,	e 202,	e 17,	161, A	e 1412	658,	e 29,	14,	14,	7, 1	e 33,	e 25,	e 34,	12,	12,	e 31,	equence 11,	11, /	equence 32,	e 30,	e 23,	13,	13, 1	e 59,	e 59,	32,	e 8,	e 28,	Sequence 15, Appl

ALIGNMENTS

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CLASSIFICATION: 536 ATTORNEY/AGENT INFORMATION: NAME: Pierri, Margaret A REGISTRATION NUMBER: 30,709 REFERENCE/DOCKET NUMBER: SIM-8	FILING DATE: 1-DEC-1994	CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/318,892	SOFTWARE: WordPerfect 6.1	OPERATING SYSTEM: PC-DOS/MS-DOS	COMPUTER: IBM PC compatible	MEDIUM TYPE: Floppy disk	COMPUTER READABLE FORM:	ZIP: 10020	COUNTRY: USA	STATE: N.Y.	CITY: New York	STREET: 1251 Avenue of the Americas	ADDRESSEE: c/o FISH & NEAVE	CORRESPONDENCE ADDRESS:	NUMBER OF SEQUENCES: 29	TITLE OF INVENTION: PEPTIDES	TITLE OF INVENTION: FOR HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING	TITLE OF INVENTION: HYALURONAN RECEPTOR (RHAMM = RECEPTOR	APPLICANT: Turley, Eva A	GENERAL INFORMATION:	Sequence 4, Application US/08318892	US-08-318-892-4	

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US-09-978-309A-31
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US-09-685-010-31
                   FILE REFERENCE: 033352-010

CURRENT APPLICATION NUMBER: US/09/978,309A

CURRENT FILING DATE: 2002-04-04

PRIOR APPLICATION NUMBER: US 09/685,010

PRIOR FILING DATE: 2000-10-05

PRIOR APPLICATION NUMBER: US 09/541,522

PRIOR FILING DATE: 2000-04-03

PRIOR APPLICATION NUMBER: US 60/127,457

PRIOR APPLICATION NUMBER: US 60/127,457

PRIOR PILING DATE: 1999-04-01

NUMBER OF SEQ ID NOS: 84
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 31
                                                                                                                                                                                                                                                                                                                                                             Sequence 31, Application US/09978309A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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                                                                                                                                                                                                                    APPLICANT: Cruz, Tony
APPLICANT: Pastrak, Aleksandra
APPLICANT: Pastrak, Aleksandra
APPLICANT: Turley, Eva A.
TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response to
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated by
TITLE OF INVENTION: Hyaladherin and Hyaluronans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATING TITLE OF INVENTION: CELLULAR RESPONSE TO INVERY AND OTHER PROLIFERATING CELL TITLE OF INVENTION: DISORDERS REGULATED BY HYALADHERIN AND HYALURONANS FILE REFERENCE: 910130.401C1
CURRENT APPLICATION NUMBER: US/09/685,010
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (212) 596-90 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Turley, Eva A. APPLICANT: Cruz, Tony F.
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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: 11
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Local Similarity 100.0%;
les 10; Conservative 0
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TYPE: amino acid
STRANDEDNESS: single
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FastSEQ for Windows Version 4.0
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linear
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ID NO: 4:
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Pred. No. 0.11;
Nismatches 0;
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RESULT 5
US-09-169-077-2
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                                                                                                                                                                                              US-08-318-892B-4
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/318,892B
FILING DATE: 1-DEC-1994
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Pierri, Margaret A
REGISTRATION NUMBER: 30709
REGISTRATION NUMBER: 30709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/08318892B GENERAL INFORMATION:
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LENGTH: 10
                                                                                                                                Matches
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Best Local Similarity
Matches 10; Conserv
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
                                                                                                                                                                                                             MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: SI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: TUX1ey, EVA A
TITLE OF INVENTION: HYALURONAN RECEPTOR
TITLE OF INVENTION: HYALURONAN MEDIATED
NUMBER OF SEQUENCES: 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                              Local Similarity hes 9; Conserv
                                                                                                                                                                                                                         STRANDEDNESS: SIR
                                                                                                                                                                                                                                                            TYPE: amino acid
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1251 Avenue of the Americas
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1; Mismatches
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Pred. No. 0.11;
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                                                                                                                                                              DB 7;
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                                                                                                                              <u>,</u>
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Sequence 2, Application US/09169077 GENERAL INFORMATION:

APPLICANT:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:

HUMAN HYALURONAN RECEPTOR 52

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NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn version 3.1
SEQ ID NO 6
LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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TOPOLOGY: linear;
MOLECULE TYPE: peptide
US-09-169-077-2
                                                                                                                                        US-09-424-292-15
                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Synthetic US-09-424-292-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-424-292-6
              Sequence 15, Application US/09424292
GENERAL INFORMATION:
APPLICANT: Turley, Eva A
TITLE OF INVENTION: Improved Delivery of Disease Modifiers
FILE REFERENCE: 205375
CURRENT APPLICATION UNMERE: US/09/424,292
CURRENT FILING DATE: 2000-07-06
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CURRENT APPLICATION NUMBER: US/09/424,292
CURRENT FILING DATE: 2000-07-06
PRIOR APPLICATION NUMBER: PCT/CA98/00448
PRIOR FILING DATE: 1998-05-11
PRIOR APPLICATION NUMBER: CA 2,205,771
PRIOR FILING DATE: 1997-05-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 10-APR-1996
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
PRIOR APPLICATION NUMBER: PCT/CA98/00448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Turley, Eva A
TITLE OF INVENTION: Improved Delivery of Disease Modifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/169,077
PTITTUE NATE.
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MEDIUM TYPE: Floppy disk
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les 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ENGTH:
                                                                                                                                                                                                                                                   1 KLKSQLVKRK 10
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                                                                                                                                                                                                               KLRSQLVKRK 10
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amino acid
DEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93.5%;
ilarity 90.0%;
Conservative
                                                                                                                                                                                                                                                                                                      93.5%;
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                                                                                                                                                                                                                                                                                   Score 43; DB 1
Pred. No. 0.42;
1; Mismatches
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Pred. No. 0.42;
1; Mismatches
                                                                                                                                                                                                                                                                                                                       DB 18; Length 10;
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Query Match
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Best Local Similarity
Thes 9; Conserve
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                                                                                                                                                                                                                                                                                            US-09-169-077-8
                                                                                                                                                                                                                                                                                                              RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1998-05-11
PRIOR APPLICATION NUMBER: CA 2,205,771
PRIOR FILING DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin version 3.1
SEQ ID NO 15
                                                                                                                                                                                                                                                          Sequence 8, Application US/09169077 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.1 SEQ ID NO 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/424,292
CURRENT FILING DATE: 2000-07-06
PRIOR APPLICATION NUMBER: PCT/CA98/00448
PRIOR FILING DATE: 1998-05-11
PRIOR APPLICATION NUMBER: CA 2,205,771
PRIOR PILING DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Turley, Eva A
TITLE OF INVENTION: Improved Delivery of Disease Modifiers
FILE REFERENCE: 205375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
OTHER INFORMATION: Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                               TITLE OF INVENTION: HUMAN HYALURONAN RECEPTOR
NUMBER OF SEQUENCES: 52
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/169,077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENGTH:
                              FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                           APPLICANT:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                             1 KLKSQLVKRK 10
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GB 9607441.4
                                WO PCT/CA97/00240
                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Mismatches
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Pred. No.
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Pred. No. 0.42;
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RESULT 11
US-09-685-010-59
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Sequence 59, Application US/09685010 GENERAL INFORMATION: APPLICANT: Turley, Eva A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 32, Application US/08318892B
GENERAL INFORMATION:
APPLICANT: Turley, Eva A
                                                                                                                                                                                                         Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (212) 596-9090
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Pierri, Margaret A
REGISTRATION NUMBER: 30709
REFERENCE/DOCKET NUMBER: SIM-8
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
TELEPAX: (212) 596-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 10-APR-1996
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Turley, Eva A
TITLE OF INVENTION: HYALURONAN RECEPTOR (RHAMM = RECEPTOR FOR
TITLE OF INVENTION: HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING PEPTIDES
NUMBER OF SEQUENCES: 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/318,892B
FILING DATE: 1-DEC-1994
CLASSIFICATION: 536
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                         TYPE: amino ac
STRANDEDNESS:
TOPOLOGY: line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 1251 Avenue of the Americas STATE: New York STATE: N.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 10020
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Local Similarity 80.0%;
les 8; Conservative
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                                                                                                                                                         1 KLKSQLVKR 9
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                                                                                                                          KLRSQLVKR 9
                                                                                                                                                                                       Similarity 88.88; Conservative
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amino acid
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                                                                                                                                                                                                       Score 38; DB 7;
Pred. No. 4.2e+06;
                                                                                                                                                                                       Mismatches
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US-08-318-892-13

Sequence 13, Application US/08318892

GENERAL INFORMATION:

APPLICANT: TUX1ey, EVA A

TITLE OF INVENTION: HYALURONAN RECI-
TITLE OF INVENTION: PEPTIDES

NUMBER OF SEQUENCES: 29

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATING
TITLE OF INVENTION: CELLULAR RESPONSE TO INJURY AND OTHER PROLIFERATING (
TITLE OF INVENTION: DISORDERS REGULATED BY HYALADHERIN AND HYALURONANS
TILE REFERENCE: 910130.401C1
CURRENT APPLICATION NUMBER: US/09/685,010
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 72
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 59
TYPE: PRT
ORGANISM: Homo sapien
US-09-685-010-59
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US-09-978-309A-59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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APPLICANT: Pastrak, Aleksandra
APPLICANT: Pastrak, Aleksandra
APPLICANT: Turley, Eva A.

TITLE OF INVENTION: Compositions and Methods for Treating Cellular Response to
TITLE OF INVENTION: Injury and Other Proliferating Cell Disorders Regulated by
TITLE OF INVENTION: Hyaladherin and Hyaluronans
FILE REFERENCE: 033552-010
CURRENT APPLICATION NUMBER: US/09/978,309A
CURRENT FILING DATE: 2000-04-04
PRIOR APPLICATION NUMBER: US 09/685,010
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 2000-04-03
PRIOR PRIOR APPLICATION NUMBER: US 09/541,522
PRIOR APPLICATION NUMBER: US 09/541,522
PRIOR PRIOR APPLICATION NUMBER: US 09/541,522
PRIOR FILING DATE: 2000-04-03
PRIOR FILING DATE: 2000-04-03
PRIOR FILING DATE: 1999-04-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 84
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
nes 8; Conserv
  STREET:
                    ADDRESSEE:
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E: c/o FISH & NEAVE
1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                             HYALURONAN RECEPTOR (RHAMM = RECEPTOR FOR HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING PEPTIDES
29
                                                                                                                                                                                                                                                                                                                                                                                                                     82.6%;
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Pred. No. 4.2e+06;
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Pred. No. 4.2e+06;
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New York

COMPUTER READABLE FORM: MEDIUM TYPE: Floppy

Floppy disk

10020

STATE: N COUNTRY:

N.Y.

USA

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; LENGTH: 10 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-318-892-13
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NAME: Pierri, Margaret A
REGISTRATION NUMBER: 30,709
REFERENCE/DOCKET NUMBER: SIM-8
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
TELEFAX: (212) 596-9090
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 13, Application US/08318892B GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: WOrdPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/318,892
FILING DATE: 1-DEC-1994
CLASSIFICATION: 536
                                                                           TELEPHONE: (212) 596-9000
TELEPAX: (212) 596-9090
INFORMATION FOR SEQ ID NO: 13
                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Pierri, Margaret A
REGISTRATION NUMBER: 30709
REFERENCE/DOCKET NUMBER: S:
                                 SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Turley, Eva A
TITLE OF INVENTION: HYALURONAN RECEPTOR (RHAMM = RECEPTOR FOR
TITLE OF INVENTION: HYALURONAN MEDIATED MOBILITY) AND HYALURONAN BINDING PEPTIDES
                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 1251 Av CITY: New York STATE: N.Y.
                                                                                                                                                                                                                                       APPLICATION NUMBER: US/
FILING DATE: 1-DEC-1994
CLASSIFICATION: 536
STRANDEDNESS:
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1251 Avenue of the Americas
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                                                                                                                                                                                                                                                                                       Sequence 23, Application US/09424292
GENERAL INFORMATION:
APPLICANT: Turley, Eva A
TITLE OF INVENTION: Improved Delivery of Disease Modifiers
FILE REFERENCE: 205375
CURRENT APPLICATION NUMBER: US/09/424,292
CURRENT FILING DATE: 2000-07-06
PRIOR APPLICATION NUMBER: PCT/CA98/00448
PRIOR APPLICATION NUMBER: CA 2,205,771
PRIOR APPLICATION NUMBER: CA 2,205,771
                                                                                                                                                                                                                         PRIOR FILING DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn version 3.1
SEQ ID NO 23
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Best Local Similarity 88.9%;
Matches 8; Conservative
                                                                Matches
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                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                              OTHER INFORMATION: Synthetic
                                                                                                                                                                                                             LENGTH: 10
                                                              Local Similarity
nes 8; Conserv
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Perfect score:
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2: /cgn2_6/ptodata/2/paa/US06_NEW_COMB.pep:*

3: /cgn2_6/ptodata/2/paa/US07_NEW_COMB.pep:*

4: /cgn2_6/ptodata/2/paa/US08_NEW_COMB.pep:*

5: /cgn2_6/ptodata/2/paa/US09_NEW_COMB.pep:*

6: /cgn2_6/ptodata/2/paa/US10_NEW_COMB.pep:*

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Copyright (c) 1993 - 2003 Compugen Ltd.
SUMMARIES
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FILE REFERENCE: 2060.0100001

RESULT 2 US-09-641-528-24854 ; Sequence 24854, Application US/09641528 ; GENERAL INFORMATION: ; APPLICANT: Sette, Alessandro ; APPLICANT: Southwood, Scott ; APPLICANT: Chesnut, Robert ; APPLICANT: Celis, Esteban ; APPLICANT: Grey, Howard ; TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS ; FILE REFERENCE, 2060, 0100001	Query Match 50.0%; Score 23; DB 5; Length 9; Best Local Similarity 50.0%; Pred. No. 3.3e+05; Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0; Qy 3 KSQLVKRK 10	PRIOR FILING DATE: 1999-11 NUMBER OF SEQ ID NOS: 5150 SOFTWARE: FASTSEQ for Wind SEQ ID NO 2553 LENGTH: 9 TYPE: PRT ORGANISM: Artificial Sequerature: PEATURE: OTHER INFORMATION: Peptid S-09-641-528-2553	APPLICANT: Sidney, John APPLICANT: Southwood, Scott APPLICANT: Southwood, Scott APPLICANT: Chesnut, Robert APPLICANT: Chesnut, Robert APPLICANT: Celis, Esteban APPLICANT: Celis, Esteban TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS TITLE OF INVENTION: USING PRETIDE AND NUCLEIC ACID COMPOSITIONS FILE REFERENCE: 2060.010001 CURRENT APPLICATION NUMBER: US/09/641,528 CURRENT FILING DATE: 2000-08-15 PRIOR APPLICATION NUMBER: US 60/172,705	ALIGNMENTS RESULT 1 US-09-641-528-2553 ; Sequence 2553, Application US/09641528 ; APPLICANT: Sette, Alessandro	27 22 47.8 8 5 US-09-641-528A-24492 Sequence 24492, A 28 22 47.8 8 5 US-09-641-528A-24533 Sequence 24553, A 29 24 47.8 8 5 US-09-641-528A-26751 Sequence 26751, A 30 22 47.8 8 5 US-09-641-528A-31258 Sequence 26800, A 31 22 47.8 8 5 US-09-641-528A-31258 Sequence 31258, A 32 24 47.8 8 5 US-09-641-528A-31258 Sequence 31258, A 33 22 47.8 8 5 US-09-641-528A-34284 Sequence 31258, A 34 22 47.8 9 5 US-09-641-528A-31258 Sequence 41648, A 35 22 47.8 9 5 US-09-641-528-3155 Sequence 2125, Ap 36 22 47.8 9 5 US-09-641-528-2125 Sequence 2125, Ap 37 22 47.8 9 5 US-09-641-528-2125 Sequence 2125, Ap 38 22 47.8 9 5 US-09-641-528-3374 Sequence 2125, Ap 40 22 47.8 9 5 US-09-641-528-3149 Sequence 3714, Ap 40 22 47.8 9 5 US-09-641-528-31135 Sequence 3714, Ap 41 22 47.8 9 5 US-09-641-528-11135 Sequence 10125, Ap 42 24 47.8 9 5 US-09-641-528-11349 Sequence 11355, Ap 43 22 47.8 9 5 US-09-641-528-1139 Sequence 11355, Ap 44 22 47.8 9 5 US-09-641-528-3149 Sequence 11355, Ap 45 22 47.8 9 5 US-09-641-528-1139 Sequence 11359, Ap 46 22 47.8 9 5 US-09-641-528-3149 Sequence 13135, Ap 47 22 47.8 9 5 US-09-641-528-3149 Sequence 13135, Ap 48 22 47.8 9 5 US-09-641-528-3139, Ap 49 5 US-09-641-528-3149 Sequence 2374, Ap 40 22 47.8 9 5 US-09-641-528-3149 Sequence 2374, Ap 40 22 47.8 9 5 US-09-641-528-3149 Sequence 2374, Ap 41 22 47.8 9 5 US-09-641-528-3149 Sequence 13135, Ap 42 22 47.8 9 5 US-09-641-528-3149 Sequence 2374, Ap 43 22 47.8 9 5 US-09-641-528-3149 Sequence 2374, Ap 44 22 47.8 9 5 US-09-641-528-3149 Sequence 2374, Ap 45 22 47.8 9 5 US-09-641-528-3149 Sequence 24299, A 45 22 47.8 9 5 US-09-641-528-3249 Sequence 24493, A

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US-09-641-528A-24854
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US-09-641-528A-2553
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APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
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PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51504
SOFTWARE: FABLSEQ for Windows Version 4.0
SEQ ID NO 24854
                                                                                                                                                                                                                                                                                                                                                                                    Sequence 24854, Application US/09641528A
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                                TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS TITUS OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS FILE REFERENCE: 2060.0100001
                                                                                                                                         APPLICANT:
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CURRENT APPLICATION NUMBER: US/09/641,528A
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CURRENT FILING DATE: 2000-08-15
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ORGANISM: Artificial Sequence
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Celis, Esteban
Grey, Howard
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Celis, Esteban
Grey, Howard
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; Pred. No. 3.3e
3; Mismatches
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APPLICANT: Chesnut, Robert
APPLICANT: Chesnut, Robert
APPLICANT: Clis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
TILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528
CURRENT PILING DATE: 2000-08-15
PRIOR APPLICATION UMBER: US 60/172,705
PRIOR APPLICATION UMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51504
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 18049
LENGTH: 10
APPLICANT: Chesnut, KOUGEL
APPLICANT: Celis, Esteban
APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: INUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528
THEORY STILING DATE: 2000-08-15
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GENERAL INFORMATION:
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PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51504
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 24854
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Best Local
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Best Local
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TYPE: PRT
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OTHER INFORMATION: Peptide Derived
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Southwood, So
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Pred. No.
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Pred. No. 3.3e+05;
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; PRIOR APPLICATION NUMBER: US 60/172,705; PRIOR FILING DATE: 1999-12-10; NUMBER OF SEO ID NOS: 51504; SOFTWARE: FASTSEQ for Windows Version 4.0; SEQ ID NO 24855
; LENGTH; 10
; TYPE: PRI
; ORGANISM: Artificial Sequence
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US-09-641-528A-24855
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US-09-641-528A-18049
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US-09-641-528-24855
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PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51504
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 18049
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Best Local Similarity
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APPLICANT:
APPLICANT:
             APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528A
CURRENT FILING DATE: 2000-08-15
                                                                                                                                                                              APPLICANT: Sette, Alessandro APPLICANT: Sidney, John
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CURRENT APPLICATION NUMBER: US/09/641,528A
CURRENT FILING DATE: 2000-08-15
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TYPE: PRT
ORGANISM: Artificial Sequence
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APPLICATION NUMBER: US 60/172,705
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Grey, Howard
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Pred. No. 72;
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Pred. No.
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APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
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APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
                                                                                                                                                                                                                                                                   Sequence 4864, Application US/09641528 GENERAL INFORMATION:
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SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2124
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 24855
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CURRENT APPLICATION NUMBER: US/09/641,528
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS FILE REPERENCE: 2060.010001 CURRENT APPLICATION NUMBER: US/09/641,528 CURRENT FILING DATE: 2000-08-15 PRIOR APPLICATION NUMBERS: US 60/172,705 PRIOR APPLICATION NUMBER: US 60/172,705 PRIOR PILING DATE: 1999-12-10
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ORGANISM: Artificial Sequence
FEATURE:
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ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Peptide Derived from Human Papillomavirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Peptide Derived from Human Papillomavirus
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Pred. No.
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Pred. No.
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NUMBER OF SEO ID NOS: 51504
SOFTWARE: FREUSEQ for Windows Version 4.0
SEQ ID NO 9364
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Best Local 9
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Best Local
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              TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS FILE REFERENCE: 2060.010001
CURRENT APPLICATION NUMBER: US/09/641,528
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
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CURRENT APPLICATION NUMBER: US/09/641,528
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
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SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4864
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: Chesnut, Robert
: Celis, Esteban
: Grey, Howard
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Celis, Esteban
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Pred. No. 3.3e+05;
2; Mismatches 0
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Pred. No.
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RESULT 14
US-09-641-528-24492
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11706
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APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esteban
APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
FILE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528
CURRENT APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR PILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51504
SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
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APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
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OTHER INFORMATION: Peptide Derived
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Chesnut, Robert
Celis, Esteban
Grey, Howard
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Pred. No.
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Pred. No. 3.3e+05;
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Sequence 2453, Application US/09641528
GENERAL INFORMATION:
APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Sidney, John
APPLICANT: Sidney, John
APPLICANT: Celis, Esteban
APPLICANT: Celis, USUBUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: UNING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528
CURRENT FILLING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILLING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51504
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 24553
LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
PETING PARTION: Peptide Derived from Human Papillomavirus
US-09-641-528-24553
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LENGTH: 8
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
GTHER IMFORMATION: Peptide Derived from Human Papillomavirus
US-09-641-528-24492
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Search completed: February 8, 2003, 11:07:26
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US-09-641-528-24553
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Best Local Similarity 66.7
Matches 4; Conservative
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Best Local Similarity 66.7
Matches 4; Conservative
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Pred. No. 3.3e+05;
2; Mismatches 0
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/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/backfiles1.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	₽B	ID	Description
1	23	50.0		r į	US-08-465-325-129	Sequence 129. App
N	23	50.0	10	4	-09-115-737	e 129,
ω	22	7.		N	-08-666-5	e 50, 1
4	22	7.	10	ω	US-08-865-472-50	e 50,
ហ	22	47.8	10	4	-09-650-438-	æ
6	22		10	ψ	93-12249	50,
7	21		9	_	64B-	e 29,
œ	21		9	N	US-08-460-421A-29	e 29,
9	21	45.7	10	N	US-08-467-202-2	æ
10	21	•	10	N	US-08-666-564-51	Sequence 51, Appl
11	21		10	w	US-08-865-472-51	Sequence 51, Appl
12	21	•	10	4	US-09-650-438-51	Sequence 51, Appl
13	21		10	v	PCT-US93-12245-51	æ
14	21	45.7	10	σ	456	
15	20		8	N	US-08-672-610A-3	•
16	20	٠	8	4	US-08-444-818-494	
17	20	43.5	9	N	1	4
18	20	٠	9	w	US-09-110-953-4	e 4,
19	20	43.5	9	4.	•	e 4,
20	20	•	10	۳		e 9,
21	20		10	_	US-08-315-831A-9	e 9,
22	20	•	10	N	-318-	е 9,
23	20	•	10	N	US-08-635-007-5	е 5
24	20	•	10	w		æ
25	20	43.5	10	w	-09-139-	44
26	20		10	ω	US-09-139-762A-60	Sequence 60, Appl
27	20	43.5	10	4	US-09-432-879-5	5, 2

20 43.5 10 5 PCT-US95-12509-9 19 41.3 5 6 1 US-08-815-688A-6 19 41.3 6 1 US-08-815-688A-6 19 41.3 7 2 US-08-680-326-92 19 41.3 8 1 US-08-135-733D-26 19 41.3 8 1 US-08-135-733D-26 19 41.3 9 1 US-08-335-733D-26 19 41.3 9 1 US-08-467-083-12 19 41.3 9 1 US-08-467-083-12 19 41.3 9 1 US-08-4814-417B-12 19 41.3 9 2 US-08-486-348A-12 19 41.3 9 2 US-08-468-545B-12 19 41.3 9 2 US-08-468-545B-12 19 41.3 9 3 US-08-159-339A-273 19 41.3 9 3 US-08-159-339A-273 19 41.3 9 3 US-08-166-660B-12 19 41.3 9 3 US-08-846-660B-12 19 41.3 10 2 US-08-764-640-60 19 41.3 10 2 US-08-774-660-19
10 5 50 6 51 6 51 6 51 6 51 6 51 6 51 6
5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5
PCT-US95-12509-9 5187078-16 US-08-815-688A-6 US-08-680-326-92 US-08-18-476A-66 US-08-335-733D-26 US-08-335-733D-26 US-08-335-733D-26 US-08-345-348-12 US-08-467-083-12 US-08-467-083-12 US-08-467-083-12 US-08-468-545B-12 US-08-468-545B-12 US-08-159-339A-273 US-08-159-339A-273 US-08-159-339A-273 US-08-159-339A-273 US-08-159-339A-273 US-08-159-339A-273 US-08-166-680B-12 US-08-466-680B-12 US-08-764-640-60

ALIGNMENTS

RESULT 1 US-08-465-325-129

; MOLECULE TYPE: US-08-465-325-129 COMPUTER READABLE FORM:

MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,325
FILING DATE: 05-UN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/184,462
FILING DATE: 18-UAN-94
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/891,201
FILING DATE: 18-UAN-94
PRIOR APPLICATION NUMBER: 07/891,201
FILING DATE: 01-UN-92
ATTORNEY/AGENT INFORMATION:
NAME: FORdis, Jean B
REGISTRATION NUMBER: 32,984
PRIPRENCE/DOCKET NIMBER: 05387.0021-03000 Sequence 129, Application US/08465325 Patent No. 5686563 GENERAL INFORMATION: TELEPHONE: (202) 408-400
TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: 1 TYPE: amino acids
STRANDEDNESS: single
TOPOLOGY: linear REGISTRATION NUMBER: 32,984
REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000 TITLE OF INVENTION: Biologically Active Peptides Having TITLE OF INVENTION: N-Terminal Substitutions NUMBER OF SEQUENCES: 153
CORRESPONDENCE ADDRESS: SEQUENCE CHARACTERISTICS: LENGTH: 10 amino acids STREET: 1300 I.S CITY: Washington STATE: D.C APPLICANT: Magainin Pharmaceuticals Inc. APPLICANT: 5110 Campus Drive APPLICANT: Plymouth Meeting, PA 19462 ZIP: 20005-3315 COUNTRY: ADDRESSEE: E: Dunner 1300 I. Street, N.W. Suite 700 USA Finnegan, Henderson, Farabow, peptide 129:

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TOPOLOGY: linear

HOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 129:
US-09-115-737-129
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                                                                                                                                         Query Match
Best Local Similarity
"---hes 4; Conserv:
US_08-666-564-50
; Sequence 50, Application US/08666564
                                    RESULT 3
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US-09-115-737-129
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Best Local Similarity
""" Ches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 129: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/465,330
FILING DATE: 05-JUN-1995
APPLICATION NUMBER: 08/184,462
FILING DATE: 18-JAN-94
APPLICATION NUMBER: 07/891,201
FILING DATE: 01-JUN-92
ATTORNEY/AGENT INFORMATION:
NAME: FORGLE, 01-JUN-92
ATTORNEY/AGENT NUMBER: 32,984
REFERENCE/DOCKET NUMBER: 05387.0021-06000
TELECOMMUNICATION INFORMATION:
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                                                                                                                       1 KLKSQLVKR 9
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOPTMARE: Patentin Release #1.0, Version #1.3

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/115,737
FILING DATE: 15-Jul-1998
CLASSIFFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Biologically Active Peptides With Reduced Toxicity in Animals and a Method for Preparing Same NUMBER OF SEQUENCES: 156 CORRESPONDENCE ADDRESS:
                                                                                      KLKKKFLKK 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: U.
                                                                                                                                                                                                                                                                                               STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dunner, L.L.P.
STREET: 1300 I Street, N.W. Suite 700
                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE:
                                                                                                                                                                                                                                                                                                                                     LENGTH: 10 amino acids
                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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Michael McLane
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                                                                                                                                                                                                                                                                                                                                                                                                          (202) 408-4000
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Pred. No. 2.1e+02;
3; Mismatches 2
                                                                                                                                                        Score 23; DB 4; Length 10;
Pred. No. 2.1e+02;
3; Mismatches 2; Indels
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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COMPUTER: IBM PC COMPATIBLE

SOPTMARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: U5/08/666,564

FILING DATE: 14-JUN-1996

CLASSIFICATION: 530

PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/US93/12245

FILING DATE: 15-DEC-1993

ATTORNEY/AGENT INFORMATION:

NAME: Kitchell, Barbara S.

REGISTRATION NUMBER: 33,928

REGISTRATION NUMBER: 33,928

REFERENCE/DOCKET NUMBER: UTSN:016
                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 512/474-7577
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, V
                                                                                                                                                                                                                           APPLICANT: Miller, Edmund J.
APPLICANT: Hayashi, Shinichiro
TITLE OF INVENTION: PEPTIDE TUMOR CELL GROWTH INHIBITORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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APPLICANT: Tuttle, Ronald R.
TITLE OF INVENTION: PEPTIDE INHIBITORS OF CXC INTERCRINE
TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES: 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee STREET: P.O. Box 4433
                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                               STREET: P.O. B
CITY: Houston
STATE: Texas
                                                                              COUNTRY: US
ZIP: 77210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
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ZIP: 77210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                      1 ELRSOSIK 8
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                                                                                                                                                      P.O. Box 4433
                                                                                                USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   512/474
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Hayashi, Shinichiro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   512/418-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             47.8%;
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                                                                                                                                                                       White & Durkee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 22; DB 2;
Pred. No. 3.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 10;
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US-09-650-438-50
; Sequence 50, Application US/09650438
; Patent No. 6355619
; GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
                                                                                            TELECOMUNICATION INFORMATION:
TELEPHONE: 512/418-3000
TELEPAX: 512/474-7577
INFORMATION FOR SEQ ID NO: 50:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 33,928
REFERENCE/DOCKET NUMBER: UTSN:020
TELECOMMUNICATION INFORMATION:
TELEPHONE: 512/418-3000
TELEFAX: 512/474-7577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 530
PRIOR APPLICATION DATA:
US 08/666,564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/
FILING DATE: 14-UUN-1996
ATTORNEY/AGENT INFORMATION:
NAME: Kitchell, Barbara S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 ELRSQSIK 8
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                                                                                                                                                                       APPLICATION NUMBER: 08/865,472
FILING DATE: -Unknown-
ATTORNEY/AGENT INFORMATION:
NAME: Kitchell, Barbara S.
REGISTRATION NUMBER: 33,928
REFERENCE/DOCKET NUMBER: UTSN:020
                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/650,438
FILING DATE: 29-Aug-2000
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Miller, Edmund J.
Hayashi, Shinichiro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: PEPTIDE TUMOR CELL GROWTH INHIBITORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: Texas
                                    TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: P.O. Box 4433
                    STRANDEDNESS: <Unknown>
                                                     LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
COPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10 amino acids
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50.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US/08/865,472
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Pred. No. 3.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
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                                                           US-08-459-064B-29
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                                                                            RESULT 7
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Sequence 29, Application US/08459064B Patent No. 5747452.
GENERAL INFORMATION:
APPLICANT: RUOSLAHTI, ERKKI I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 50, Application PC/TUS9312245 GENERAL INFORMATION:
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                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                          TELEFAX: (713) 789-26:
TELEX: 79-0924
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: UTFN016PCT TELECOMMUNICATION INFORMATION: TELEPHONE: (512) 320-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: KITCHELL, Barbara S.
REGISTRATION NUMBER: 33,91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: PC
FILING DATE: Concurrent
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: HAYASHI, Shinichiro
APPLICANT: TUTTLE, Ronald R.
TITLE OF INVENTION: PEPTIDE INHIBITORS OF CXC
TITLE OF INVENTION: INTERCRINE MOLECULES
                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                     Local Similarity tes 4; Conserv
                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
SOFTWARE: WordPerfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: P. O. CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
les 4; Conserv
                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                           TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 77210
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                                                                                                                         1 ELRSQSIK 8
                                                                                                                                                        1 KLKSQLVK 8
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                                                                                                                                                                                                                                                                                                                               10 amino acids
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                                                                                                                                                                                                                                                                             linear
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HAYASHI, Shinichiro
                                                                                                                                                                                     47.8%;
llarity 50.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   UMBER: PCT/US93/12245
Concurrently herewith
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50.0%;
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                                                                                                                                                                                     Score 22; DB Pred. No. 3.1e 3; Mismatches
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Pred. No.
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3.1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 45.7%;
Best Local Similarity 71.4%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -08-459-064B-29
                                                                                                                                                                                                                                                                                                                                        Patent No. 5837813
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/829,462
FILING DATE: 31-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/021,626
FILING DATE: 16-FEB-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/340,812
APPLICATION NUMBER: US 08/340,812
PILING DATE: 17-NOV-1994
ATTORNEY/ACENT INFORMATION:
NAME: CAMPBERI CATTLEYN
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 619-535-8949
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                        APPLICANT: RUOSLAHTI, APPLICANT: MORLA, ALEX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 619-535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/459,064B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: A METHOD OF MODULATING TUMOR CELL MIGRATION TITLE OF INVENTION: USING FIBRONECTIN TYPE III PEPTIDES NUMBER OF SEQUENCES: 35
                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: CAMPBELL, CATHRYN A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LA 1543
                                                                                                                                      STREET: 4370 LA
CITY: SAN DIEGO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 01-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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                                                                                           COUNTRY: UNITED STATES
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                                                                                                                  CALIFORNIA
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4370 LA JOLLA VILLAGE DRIVE, STE 700
                                                                                                                                                        3: CAMPBELL & FLORES LLP
4370 LA JOLLA VILLAGE DRIVE,
                                                                                                                                                                                                                                                                                                                 RUOSLAHTI, ERKKI I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein
                                                                                                                                                                                                                                               FIBRONECTIN BINDING SITES AND METHODS OF MODULATING FIBRONECTIN EXTRACELLULAR MATRIX ASSEMBLY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 21; DB 1; Length 9; Pred. No. 2e+05; 0; Mismatches 2; Indels
                                                                                                                                                           STE 700
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Query Match
Best Local Similarity
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Patent No. 5840295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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FILING DATE: 31-JAX-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 16-FEB-1993
PRIOR APPLICATION DATA: US 08/
PRIOR APPLICATION NUMBER: US 08/
APPLICATION NUMBER: US 08/
FILING DATE: 17-NOV-1994
ATTORNEY/AGENT INFORMATION:
NAME: CAMPBELL, CATHRYN A.
REGISTRATION NUMBER: 31,81
REGESTRATION NUMBER: 31,81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 06
PILING DATE: 30-JUL-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                   COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                ZIP: 60606-0110
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Enzymatically-Produced Dimeric IL-2 and A TITLE OF INVENTION: Nerve-Derived Transglutaminase Enzyme for Its Preparation NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,421A
FILING DATE: 01-JUN-1995
                APPLICATION NUMBER: US 07/840,783
FILING DATE: 24-FEB-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Schwartz, Michal APPLICANT: Eitan, Shoshana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: 619-535-9001
                                                                                                                                                         APPLICATION NUMBER: US/08/467,202 FILING DATE: 07-JUN-1995 CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
APPLICATION NUMBER:
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GY: linear
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P.O. Box 06110
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71.4%;
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                                                                                                                    US 08/099,759
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  US 07/573,580
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Pred. No. 2e+05;
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 30-JUL-

30-JUL-1992

IL 102686

DATE:

27-AUG-1990

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RESULT 10
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TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-467-202-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 51, Approx No. 5965536
COMFUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/666,564
FILING DATE: 14-JUN-1996
CLASSIFICATION 1530
PRIOR APPLICATION NUMBER: PCT/US93/12245
APPLICATION NUMBER: PCT/US93/12245
APPLICATION NUMBER: PCT/US93/12245
APPLICATION NUMBER: DCT/US93/12245
APPLICATION NUMBER: US/08/12245
APPLICATION NUMBER: DCT/US93/12245
APPLICATION NUMBER: DCT/US93/12245
FILING DATE: 15-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: Kitchell, Barbara S.
REGISTRATION NUMBER: UTSN:016
TELEPHONE: 512/418-3000
TELEPHONE: 512/418-3000
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TELEFAX: (202) 789-1158
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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NAME: WALBON T. SCOUL
REGISTRATION NUMBER: 26,581
REFERENCE/DOCKET NUMBER: 47142-053
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Kurdowska, Anna
APPLICANT: Hayashi, Shinichiro
APPLICANT: Tuttle, Ronald R.
TITLE OF INVENTION: PEPTIDE INHIBITORS OF CXC INTERCRINE
TITLE OF INVENTION: MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: F. STREET: F. STREET: Fouston
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: IL 1:
FILING DATE: 19-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: US
ZIP: 77210
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Local Similarity 57.1%;
1es 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Arnold, White & Durkee P.O. Box 4433
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Miller, Edmund J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cohen, Allen B.
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Best Local Similarity
Matches 3; Conserv
                                                                                                                                                                                                                                                                            COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/865,472
FILING DATE: CONCURRENTLY HEREWITH
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/666,564
FILING DATE: 14-UN-1996
ATTORNEY/AGENT INFORMATION:
NAME: Kitchell, Barbara'S.
REGISTRATION NUMBER: 33,928
REFERENCE/DOCKET NUMBER: UTSN:020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 6110889
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 51, Application US/08865472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                 TELEFAX: 512/474-7577
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Miller, Edmund J.
APPLICANT: Hayashi, Shinichiro
TITLE OF INVENTION: PEPTIDE TUMOR CELL GROWTH INHIBITORS
NUMBER OF SEQUENCES: 58
CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee
STREET: P.O. Box 4433
                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
                                                                                                                                 COUNTRY:
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1 ELRMQMIK 8
                           1 KLKSQLVK 8
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; Pred. No. 4.7e
4; Mismatches
                                                          Score 21; DB
Pred. No. 4.7e
4; Mismatches
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US-09-650-438-51 ; Sequence 51, Application US/09650438 ; Patent No. 6355619

RESULT 12

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NAME: Kitchell, Barbara S.

REGISTRATION NUMBER: 33,928

REFERENCE/DOCKET NUMBER: UTSN:020

TELECOMMUNICATION INFORMATION:

TELEPHONE: 512/418-3000

TELEPHONE: 512/418-3000

TELEPAX: 512/418-7577

INFORMATION FOR SEQ ID NO: 51:

SEQUENCE CHARACTERISTICS:

SEQUENCE CHARACTERISTICS:

TYPE: amino acids

TYPE: amino acids

TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 51, Applicat GENERAL INFORMATION:
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APPLICANT: Miller,
ZIP: 77210
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
                                                                                                                                                                                                                                              APPLICANT: COHEN, Allen B.
APPLICANT: MILLER, Edmund J.
APPLICANT: KURDOWSKA, Anna
APPLICANT: HAYASHI, Shinichiro
APPLICANT: TUTTLE, Ronald R.
TITLE OF INVENTION: PEPTIDE INHIBITORS OF CXC
TITLE OF INVENTION: INTERCRINE MOLECULES
                                                                                                                                                                                                          NUMBER OF SEQUENCES: 5
                                                                                                                          STREET: ..
STREET: F.
TTY: Houston
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les 3; Conserv
                                                                                                     COUNTRY:
                                                                                                                                                                                       ADDRESSEE:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08/865,472 FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/650,438
FILING DATE: 29-Aug-2000
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Miller, Edmund J.
TITLE OF INVENTION: PEPTIDE TUMOR CELL GROWTH INHIBITORS
NUMBER OF SEQUENCES: 58
CORRESPONDENCE ADDRESS:
ADDRESSEE: Arrold, White & Durkee
STREET: P.O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                       Texas
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                                                                                                                                                                  E: Arnold, White & Durkee P. O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                      Application PC/TUS9312245
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37.5%;
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Pred. No. 4.7e+02;
4; Mismatches 1; Indels
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5514565-10.
5514565-10.
5514565-10.
APPLICANT: SCHWARTZ, MICHAEL; EITAN, SHOSHANA
TITLS OF INVENTION: ENZYMATICALLY-PRODUCED DIMERIC IL-2;
AND A NERVE-DERIVED TRANSGLUTAMINASE ENZYME FOR ITS PREPARATION
OMBER OF SEQUENCES: 10
CURRENT APPLICATION UNMBER: US/08/99,759
FILING DATE: 30-JUL-1993
PRIOR APPLICATION UNMBER: US/08/99,759
FILING DATE: 30-JUL-1993
PRIOR APPLICATION DATA:
THE PRIOR APPLICATION DATA:
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THE PRIOR APPLICATION DATA:
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                                                                                                                                                                               Sequence 3, Application US/08672610A Patent No. 5846933 GENERAL INFORMATION:
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Matches 3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KEGISTRATION NUMBER: 33,928
REFERENCE/DOCKET NUMBER: UTFN016PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 320-7200
TELEPAX: (713) 789-2679
TELEPAX: (713) 789-2679
                                                           NUMBER OF SEQUENCES: 5
                                                                                                 APPLICANT: Ziwei, Hwang
APPLICANT: Korngold, Robert
TITLE OF INVENTION: CD4-DERIVED PEPTIDES THAT INHIBIT IMMUNE
TITLE OF INVENTION: RESPONSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDENESS: single
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APPLICATION NUMBER: PCT/US93/12245
FILING DATE: Concurrently herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                       ADDRESSEE:
STREET: 11
                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
nes 4; Conserv
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FILING DATE: 24-FEB-1992
APPLICATION NUMBER: 573,580
FILING DATE: 27-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: KITCHELL, Barbara S. REGISTRATION NUMBER: 33,92
                                                                                                                                                                                                                                                                                                                               2 SKLTKKK 8
                                                                                                                                                                                                                                                                                                                                                                     4 SQLVKRK 10
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    New York
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                     E: PENNIE & EDMONDS
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; Pred. No. 4.7e.
2; Mismatches
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Length 10;

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Gaps

0;

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Score 21; DB 5;
Pred. No. 4.7e+02;
 Mismatches
                  Length 10
<u>,,</u>
Gaps
0
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COMPTRY: United States
21P: 10036-2711
COMPUTER RADABLE FORM:
KMEDIUM TYPE: Floppy disk
COMPUTER: EADABLE FORM:
COMPUTER: EADABLE FORM:
COMPUTER: EADABLE FORM:
COMPUTER: IBM PC COMPATAIN:
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